

(SEQ. ID. NO. 195)

MEGLVRVHLLPVFGDYKLSKLTTPILQQQVNKWADKANKGEKGAFANYSLLHNMNKRILKYGVAIQVIQYNPANDVIVPRKQQ KEKAAVKYLDNKELKQFLDYLDALDQSNYENLFDVVLYKTLLATGCRISEALALEWSDIDLESGVISINKTLNRYQEINSPKS SAGYRDIPIDKATLLLLKQYKNRQQIQSWKLGRSETVVFSVFTEKYAYACNLRKRLNKHFDAAGVTNVSFHGFRHTHTTMMLY AQVSPKDVQYRLGHSNLMITENTYWHTNQENAKKAVSNYETAINNLZ--

Please delete the first paragraph located on page 101, line 1, and replace therewith the following replacement paragraph. A marked-up version showing changes made to the replacement paragraphs is contained herewith as Appendix B.



--What is claimed is:--

REMARKS

Claims 1-20 are currently pending in this application. Upon entry of the amendments contained herein, claims 1-22 will be pending. No new matter is introduced into the specification by way of these amendments. The Specification amendments are introduced to conform the application to the requirements of 37 C.F.R. § § 1.77 and 1.821-1.825. Support for the claim amendments can be found, inter alia, throughout the application as originally filed, for example, in the originally filed claims; on page 9, last two full paragraphs; and in Table 3. Support for new claim 21 can be found, inter alia, throughout the specification, for example, on page 9, last two full paragraphs. Support for new claim 22 can be found, inter alia, throughout the Specification, for example, on page 9, last two full paragraphs and on page 13, third full paragraph.

Priority

The Examiner has made acknowledgement of Applicant's claim of foreign priority based on "German" foreign application number 9816337.1. Applicants respectfully state that the claim to foreign priority is in fact based upon Great Britain priority application number 9816337.1. Applicants respectfully direct the Examiner's attention to the executed declaration, claiming the benefit of GB 9816337.1, as well as the cover sheet of International Publication Number WO 00/06738 (reciting the priority application GB 9816337.1, and of which the instant application claims benefit), attached herewith as Appendix C. Applicants will provide the Examiner with a certified copy of Great Britain priority application GB 9816337.1 within a reasonable time of receipt of the document.

Restriction Requirement

In the Office Action dated September 22, 2002, the Examiner required restriction of the pending claims under 35 U.S.C. § 121 between one of the following inventions:

- "1.) Claims 7-8 and 13 drawn to DNA, host cell and a vaccine composition, classified in class 536, subclass 23.7. Further election of invention required.
- 2.) Claims 1-6 and 10-12 drawn to polypeptide, classified in class 424, subclass 190.1. Further election of invention required.
- 3.) Claims 15-16 drawn to an antibody classified in class 530, subclass 388.6. Further election of invention required.
- **4.)** Claim 9 drawn to a method for inducing immune response using polypeptide classified in class 424, subclass 165.1. Further election of invention required.
- 5.) Claim 20 drawn to a method of use for the treatment or prophylaxis of S. pneumoniae infection using protein or polypeptide classified in class 424, subclass 184.1. Further election of invention required.
- **6.)** Claim 19 drawn to a method of determining whether a protein is anti-microbial target in vivo or in vitro classified in class 424, subclass 93.1. Further election of invention required.
- 7.) Claim 18 drawn to a method of detecting S. pneumoniae infection using nucleic acid, SEQ ID NO:1 classified in class 435, subclass 6. Further election of invention required.
- **8.)** Claim 14 drawn to a method of detecting S. pneumoniae infection using protein antigen, SEQ ID NO:2, classified in class 435, subclass 7.22. Further election of invention required.
- 9.) Claim 17 drawn to a method of detecting S. pneumoniae infection using antibody classified in class 435, subclass 7.22. Further election of invention required."

See Paper No. 11, page 2. The Examiner further required election of one sequence falling within one of the above-identified inventions.

Applicants were contacted by the Examiner and invited to make an election to a restriction requirement by telephone. The Examiner originally designated (by telephone) as separate inventions the following 5 groups: (I) *S. pneumoniae* nucleic acid sequences (claims 7-8 and 18); (II) *S. pneumoniae* peptide sequences (claims 1-6 and 9-14); (III) antibodies to *S. pneumoniae* antigens (claims 15-17); (IV) methods of assay design (claim 19); and therapeutic methods (claim 20). Based upon this grouping of claims, Applicants provisionally elected by telephone on July 09, 2002, without traverse, the invention of Group V, claim 20, *drawn to therapeutic methods*. To the extent that the restriction requirement in the Office Action dated September 20, 2002, differs from that described in the telephone restriction to which Applicants responded on July 9, 2002, the restriction is traversed. In particular, Applicants submit that there is substantial overlap in scope between Group 4 and Group 5 or Group 6. Dependent claims to particular embodiments may properly fall within the scope of both generic claim 9 and generic claim 20. Likewise, dependent claims to other

embodiments may properly fall within the scope of both generic claim 9 and generic claim 19. Applicants respectfully request that the restriction requirement be recast.

Applicants also submit that the requirement for election of a single nucleotide (polypeptide) sequence is properly characterized as an election of species. Applicants provisionally elect the invention of claim 20, now classified in Group 5.

Consistent with the requirement of election of species, Applicants hereby further elect the protein sequence as set forth in SEQ ID NO:162.

Rejections

35 U.S.C. § 101

The Examiner rejected claim 20 under 35 U.S.C. § 101, as allegedly improperly defining a process. Applicants have amended claim 20 to conform with U.S. Rules of Practice, thereby rendering the Examiner's rejection moot. Accordingly, Applicants respectfully request reconsideration and withdrawal of the rejection of claim 20 under 35 U.S.C. § 101.

35 U.S.C. § 112, 2nd Paragraph

The Examiner rejected claim 20 under 35 U.S.C. § 112, 2nd Paragraph as allegedly indefinite for failing to particularly point out and distinctly claim the subject matter which Applicants regard as the invention. More specifically, the Examiner stated that "the claim does not set forth any steps involved in the method/process, [and therefore] it is unclear what method/process applicant is intending to encompass." See Paper No. 11, Page 7, lines 4-6. Similarly, the Examiner stated that the claim is further indefinite because "the claims do not recite positive active steps so that the claims will set out and circumscribe particular area with reasonable degree of precision and particularly [sic] and make clear what subject matter the claims encompass, as well as make clear the subject matter from which others would be precluded." See Paper No. 11, Page 7, lines 9-13.

The Examiner further determined that "additional structural parameters, such as amino acid sequence of the polypeptide (for example; SEQ.ID.NO), would enable more definitive identification of the claimed polypeptide without ambiguity." See Paper No. 11, Page 7, lines 17-19.

The Examiner also rejected claim 20 as allegedly vague for the recitation of "capable of" within the claim, as this recitation allegedly makes it difficult to determine the metes and bounds of the claim.

Applicants have amended claim 20 to confirm to U.S. practice. As a result of the claim amendment, the Examiner's rejection of claim 20 has been rendered moot. Applicants respectfully request withdrawal of the rejection of claim 20 under 35 U.S.C. § 112, 2nd Paragraph, and reconsideration of the application in view of the amendments.

Conclusion

Applicants believe that amendment of the subject application has placed this application in a condition for allowance. Early notification of a favorable consideration is respectfully requested.

Respectfully submitted,

Dated: March 20, 2003

By:

Laurence H. Posorske Registration No. 34,698 Robert C. Lampe III Registration No. 51,914

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Facsimile: (202) 778-2201

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re P	atent Application of:)	
	GILBERT, C.F.G., et al.)) E	xaminer: Baskar, P.
Serial	No.: 09/769,787)) G	roup Art Unit: 1645
Filed:	January 26, 2001)	
For:	NUCLEIC ACIDS AND PROTEINS FROM STREPTOCOCCUS PNEUMONIAE		

Under Secretary of Commerce for Intellectual Property and Director of the United States Patent and Trademark Office

Washington, D.C. 20231

Appendix A - Version with Markings to show changes made to the Claims

In claim 20:

--20. (Amended) A method for treatment or prophylaxis of S. pneumoniae infection comprising administering to a patient in need thereof The use of an agent capable of antagonizing, inhibiting or otherwise interfering with the function or expression of a protein or polypeptide as defined in Tables 2-4 in the manufacture of a medicament for use in the treatment or prophylaxis of S. pneumoniae infection, wherein said agent induces an immune response in said patient to a polypeptide having a sequence comprising SEQ. ID NO:162.--



In re P	atent Application of:)	
	GILBERT, C.F.G., et al.	Examiner: Baskar, P.	
Serial	No.: 09/769,787) Group Art Unit: 1645	
Filed:	January 26, 2001)	
For:	NUCLEIC ACIDS AND PROTEINS FROM STREPTOCOCCUS PNEUMONIAE		
	Secretary of Commerce for Intellectual or of the United States Patent and Trade	- ·	

Appendix B - Version with Markings to show changes made

In the paragraphs starting at page 29, 1st full paragraph (immediately below the heading "Table 1") and extending to and including the last full paragraph located on page 100:

4101.1 (SEQ. ID. NO. 208)

Washington, D.C. 20231

4101.3 (SEQ. ID. NO. 209)

ATGGGGAAAGGCCATTGGAATCGGAAAAGAGTTTATAGCATTCGTAAGTTTGCTGTGGGAGCTTGCTCAGTAATGATTGGGA CTTGTGCAGTTTTATTAGGAGGAAATATAGCTGGAGAATCTGTAGTTTATGCGGATGAAACACTTATTACTCATACTGCTGAG AAACCTAAAGAGGAAAAAATGATAGTAGAAGAAAAAGGCTGATAAAGCTTTGGAAACTAAAAATATAGTTGAAAGGACAGA ACAAAGTGAACCTAGTTCAACTGAGGCTATTGCATCTGAGAAGAAGAAGATGAAGCCGTAACTCCAAAAGAGGAAAAAGT GTCTGCTAAACCGGAAGAAAAGCTCCAAGGATAGAATCACAAGCTTCAAATCAAGAAAAACCGCTCAAGGAAGATGCTAA AGCTGTAACAAATGAAGAAGTGAATCAAATGATTGAAGACAGGAAAGTGGATTTTAATCAAAATTGGTACTTTAAACTCAAT GCAAATTCTAAGGAAGCCATTAAACCTGATGCAGACGTATCTACGTGGAAAAAATTAGATTTACCGTATGACTGGAGTATCT TTAACGATTTCGATCATGAATCTCCTGCACAAAATGAAGGTGGACAGCTCAACGGTGGGGAAGCTTGGTATCGCAAGACTTT CAAACTAGATGAAAAAGACCTCAAGAAAAATGTTCGCCTTACTTTTGATGGCGTCTACATGGATTCTCAAGTTTATGTCAATG GTCAGTTAGTGGGGCATTATCCAAATGGTTATAACCAGTTCTCATATGATATCACCAAATACCTTCAAAAAGATGGTCGTGA TTACAAGTGACAGATAAGGTGCATGTTGAGAAAAATGGGACAACTATTTTAACACCAAAAACTTGAAGAACAACAACATGGC AAGGTTGAAACTCATGTGACCAGCAAAATCGTCAATACGGACGACAAAGACCATGAACTTGTAGCCGAATATCAAATCGTTG AACGAGGTGGTCATGCTGTAACAGGCTTAGTTCGTACAGCGAGTCGTACCTTAAAAGCACATGAATCAACAAGCCTAGATGC GATTTTAGAAGTTGAAAGACCAAAACTCTGGACTGTTTTAAATGACAAACCTGCCTTGTACGAATTGATTACGCGTGTTTACC GTGACGGTCAATTGGTTGATGCTAAGAAGGATTTGTTTGGTTACCGTTACTATCACTGGACTCCAAATGAAGGTTTCTCTTTG AATGGTGAACGTATTAAATTCCATGGAGTATCCTTGCACCACGACCATGGGGCGCTTGGAGCAGAAGAAACTATAAAGCA GAATATCGCCGTCTCAAACAAATGAAGGAGATGGGAGTTAACTCCATCCGTACAACCCACAACCCTGCTAGTGAGCAAACCT TGCAAATCGCAGCAGAACTAGGTTTACTCGTTCAGGAAGAGGCCTTTGATACGTGGTATGGTGGCAAGAAACCTTATGACTA TGGACGTTTCTTTGAAAAAGATGCCACTCACCCAGAAGCTCGAAAAAGGTGAAAAATGGTCTGATTTTGACCTACGTACCATG



GTCGAAAGAGCCAAAAACAACCCTGCTATCTTCATGTGGTCAATTGGTAATAGAAATAGGTGAAGCTAATGGTGATGCCCACT CTTTAGCAACTGTTAAACGTTTGGTTAAGGTTATCAAGGATGTTGATAAGACTCGCTATGTTACCATGGGAGCAGATAAATTC CGTTTCGGTAATGGTAGCGGAGGGCATGAGAAAATTGCTGATGAACTCGATGCTGTTGGATTTAACTATTCTGAAGATAATT ACAAAGCCCTTAGAGCTAAGCATCCAAAATGGTTGATTTATGGATCAGAAACATCTTCAGCTACCCGTACACGTGGAAGTTA CTATCGCCCTGAACGTGAATTGAAACATAGCAATGGACCTGAGCGTAATTATGAACAGTCAGATTATGGAAATGATCGTGTG GGTTGGGGGAAAACAGCAACCGCTTCATGGACTTTTGACCGTGACAACGCTGGCTATGCTGGACAGTTTATCTGGACAGGTA CGGACTATATTGGTGAACCTACACCATGGCACAACCAAAATCAAACTCCTGTTAAGAGCTCTTACTTTGGTATCGTAGATACA GCCGGCATTCCAAAACATGACTTCTATCTCTACCAAAGCCAATGGGTTTCTGTTAAGAAGAAACCGATGGTACACCTTCTTCC TCACTGGAACTGGGAAAACAAAGAATTAGCATCCAAAGTAGCTGACTCAGAAGGTAAGATTCCAGTTCGTGCTTATTCGAAT CTTACCAAGAAGGTGCAAATGCTAATGAACTTTATCTTGAATGGAAAGTTGCCTATCAACCAGGTACCTTGGAAGCAATTGC TCGTGATGAATCTGGCAAGGAAATTGCTCGAGATAAGATTACGACTGCTGGTAAGCCAGCGGCAGTTCGTCTTATTAAGGAA GACCATGCGATTGCAGCAGATGGAAAAGACTTGACTTACATCTACTATGAAATTGTTGACAGCCAGGGGAATGTGGTTCCAA ACGCTATAAGGCGCAAGCAGATGGTTCTTGGATTCGTAAAGCATTTAATGGTAAAGGTGTTGCCATTGTCAAATCAACTGAA CAAGCAGGGAAATTCACCCTGACTGCCCACTCTGATCTCTTGAAATCGAACCAAGTCACTGTCTTTTACTGGTAAGAAGAAG GACAAGAGAAGACTGTTTTGGGGACAGAAGTGCCAAAAGTACAGACCATTATTGGAGAGGCACCTGAAATGCCTACCACTG TTCCGTTTGTATACAGTGATGGTAGCCGTGCAGAACGTCCTGTAACCTGGTCTTCAGTAGATGTGAGCAAGCCTGGTATTGTA ACGGTGAAAGGTATGGCTGACGGACGAGAAGTAGAAGCTCGTGTAGAAGTGATTGCTCTTAAAATCAGAGCTACCAGTTGTG TGAAGTGGACAAGTGGGAGATTGCCGAAGAAGATAAAGCTAAGTTAGCAATTCCAGGTTCTCGTATTCAAGCGACCGGTTAT TTAGAAGGTCAACCAATTCATGCAACCCTTGTGGTAGAAGAAGGCAATCCTGCGGCACCTGCAGTACCAACTGTAACGGTTG GTGGTGAGGCAGTAACAGGTCTTACTAGTCAAAAACCAATGCAATACCGCACTCTTGCTTATGGAGCTAAGTTGCCAGAAGT CCTAAAGATGGTGGCCCTCTTCAAACCTATGCAATTCAATTCCTTGAAGAAGCGCCAAAAATTGCTCACTTGAGCTTGCAAGT GGAAAAAGCTGACAGTCTCAAAGAAGACCAAACTGTCAAATTGTCGGTTCGAGCTCACTATCAAGATGGAACGCAAGCTGT ATTACCAGCTGATAAAGTAACCTTCTCTACAAGTGGTGAAGGGGAAGTCGCAATTCGTAAAGGAATGCTTGAGTTGCATAAG CCAGGAGCAGTCACTCTGAACGCTGAATATGAGGGAGCTAAAGACCAAGTTGAACTCACTATCCAAGCCAATACTGAGAAG AAGATTGCGCAATCCATCCGTCCTGTAAATGTAGTGACAGATTTGCATCAGGAACCAAGTCTTCCAGCAACAGTAACAGTTG TGAAGTACTAGGTAAAGTTGAAGGAATTGACCTTGAAGCGCGTGCAAAAGTCTCTGTAGAAGGTATCGTTTCAGTTGAAGAA GTCAGTGTGACAACTCCAATCGCAGAAGCACCACAATTACCAGAAAGTGTTCGGACATATGATTCAAATGGTCACGTTTCAT TACGCAATTAACAACTAAACTTCATGTTCGCGTATCTGCTCAAACTGAGCAAGGTGCAAACATTTCTGACCAATGGACCGGTT CAGAATTGCCACTTGCCTTTGCTTCAGACTCAAATCCAAGCGACCCAGTTTCAAATGTTAATGACAAGCTCATTTCCTACAAT AACCAACCAGCCAATCGTTGGACAAACTGGAATCGTACTAATCCAGAAGCTTCAGTCGGTGTTCTGTTTGGAGATTCAGGTA TCTTGAGCAAACGCTCCGTTGATAATCTAAGTGTCGGATTCCATGAAGACCATGGAGTTGGTGTACCGAAGTCTTATGTGATT GAGTATTATGTTGGTAAGACTGTCCCAACAGCTCCTAAAAACCCTAGTTTTGTTGGTAATGAGGACCATGTCTTTAATGATTC TGCCAACTGGAAACCAGTTACTAATCTAAAAGCCCCTGCTCAACTCAAGGCTGGAGAAATGAACCACTTTAGCTTTGATAAA GTTGAAACCTATGCTGTTCGTATTCGCATGGTTAAAGCAGATAACAAGCGTGGAACGTCTATCACAGAGGTACAAATCTTTG CGAAACAAGTTGCGGCAGCCAAGCAAGGACAAACAAGAATCCAAGTTGACGGCAAAGACTTAGCAAACTTCAACCCTGATT TGACAGACTACTACCTTGAGTCTGTAGATGGAAAAGTTCCGGCAGTCACAGCAAGTGTTAGCAACAATGGTCTCGCTACCGT CGTTCCAAGCGTTCGTGAAGGTGAGCCAGTTCGTGTCATCGCGAAAGCTGAAAATGGCGACATCTTAGGAGAATACCGTCTG CACTTCACTAAGGATAAGAGCTTACTTTCTCATAAACCAGTTGCTGCGGTTAAACAAGCTCGCTTGCTACAAGTAGGTCAAGC ACTTGAATTGCCGACTAAGGTTCCAGTTTACTTCACAGGTAAAGACGGCTACGAAACAAAAGACCTGACAGTTGAATGGGAA GAAGTTCCAGCGGAAAATCTGACAAAAGCAGGTCAATTTACTGTTCGAGGCCGTGTCCTTGGTAGTAACCTTGTTGCTGAGA TCACTGTACGAGTGACAGACAAACTTGGTGAGACTCTTTCAGATAACCCTAACTATGATGAAAACAGTAACCAGGCCTTTGC TTCAGCAACCAATGATATTGACAAAAACTCTCATGACCGCGTTGACTATCTCAATGACGGAGATCATTCAGAAAATCGTCGT TGGACAAACTGGTCACCAACACCATCTTCTAATCCAGAAGTATCAGCGGGTGTGATTTTTCCGTGAAAATGGTAAGATTGTAG AACGGACTGTTACACAAGGAAAAGTTCAGTTCTTTGCAGATAGTGGTACGGATGCACCATCTAAACTCGTTTTAGAACGCTA AATTGGGAAGCTGTTCCTTATCGTGCGGATAAAGACATTGCAGCTGGTGATGAAATCAACGTAACATTTAAAGCTATCAAAG TGAATTGCCTCAAGAAAGCACTCAATCAAAGATTCTTGTAGATGGAAAAGAACTTGCTGATTTCGCTGAAAATCGTCAAGAC TATCAAATTACCTATAAAGGTCAACGGCCAAAAGTCTCAGTTGAAGAAAACAATCAAGTAGCTTCAACTGTGGTAGATAGTG GAGAAGATAGCTTTCCAGTACTTGTTCGCCTCGTTTCAGAAAGTGGAAAACAAGTCAAGGAATACCGTATCCACTTGACTAA

4101.5 (SEQ. ID. NO. 210)



4102.1 (SEQ. ID. NO. 211)

GTAATTTATCAGATTTTAGCAATAAAGGGGATTCTGATGCAATATATGCGCCTGCAGGCTCAACATTATCTCTTTCAGAATTA GGACTTAATAACTTTATTAATGCAGAAAAATATAAAGAAGATTGGATTTTTTTCGGCAACACTAGGAGGATATACGTATCTTT ATGGAAACTCATTTGCTGCTCCTAAAGTTTCTGGTGCGATTGCAATGATTATTGATAAAATACAAATTAAAAGATCAGCCCTAT AATTATATGTTTGTAAAAAAAATTCTGGAAGAAACATTACCAGTAA

4106.1 (SEQ. ID. NO. 212)

4106.4 (SEQ. ID. NO. 213)

AGCTATTTATTTGACCTTTTCATTATTATTCAACTGGAGGGGAAAAGTTTAATGAGAGCGCAAGAGTGTTTACGGAGT ATTTAAAGACTAAGACATCTGATGAAAATTCCAAGCTTACTCCAGTCTTATTCAAAGTCCTTGACCATATCTGCTCACCTTAAA AGAGATATTGTAGATAAGCGGCTCCCTCTTGTGCATGACTTGGATATTAAAGATGGAAAGCTATCAAATTATATCGTGATGTT AATATTTTGCTTTTGTATCTCCCATATACATTTTTGGTTACAATTGCTTTTTCCTTTGTTTTTTCTTATTTTTATACTAAACGCTT GCTCAATCCTCTTTTTTACATTTCAGAAGTGACTAGTAAAATGCAAGATTTTGGATGACAATATTCGTTTTGATGAAAGTAGGA TAATGAGCAAATTGTAAAATTGCAAAATCAAAAGGTTTCCTTTGTCCGCGGAGCATCACATGAGTTGAAAACCCCTTTAGCC AGTCTTAGAATTATCCTAGAGAATATGCAGCATAATATTGGAGATTACAAAGATCATCCAAAATATATTGCAAAGAGTATAA ATAAGATTGACCAGATGAGCCACTTATTAGAAGAAGTACTGGAGTCTTCTAAATTCCAAGAGTGGACAGAGTGTCGTGAGAC CTTGACTGTTAAGCCAGTTTTAGTAGATATTTTATCACGTTATCAAGAATTAGCTCATTCAATAGGTGTTACAATTGAAAATC AATTGACAGATGCTACCAGGGTCGTCATGAGTCTTAGGGCATTGGATAAGGTTTTGACAAACCTGATTAGTAATGCAATTAA ATATTCAGATAAAAATGGGCGTGTAATCATATCCGAGCAAGATGGCTATCTCTCTATCAAAAATACATGTGCGCCTCTAAGT GACCAAGAACTAGAACATTTATTTGATATATTCTATCATTCTCAAATCGTGACAGATAAGGATGAAAGTTCCGGTTTGGGTCT TTACATTGTGAATAATATTTTAGAAAGCTATCAAATGGATTATAGTTTTCTCCCTTATGAACACGGTATGGAATTTAAGATTA **GCTTGTAG**

4106.6 (SEQ. ID. NO. 214)

4106.7 (SEQ. ID. NO. 215)

TCCATTGATTGCAGGGAACAGTGCCTCTGACTATAATGGTGGAAATAACGGTACCATCAGTGATGAAAAGCTTTGACAACTTG
ATTGCGACTGTTGAATCTTATCTCTCCAAAGAAAAAACACGTGAAGATGTTGAGTCTGTCAGCAAGCTTGAAAGTAGCA
CATCTGAGAAACATTTGGATCCATCTGCAGTTTCTCGTGGGTCTAGCTTGGACCGTGATGACAATGGTCTCTTGACTCTTGCT
GGTGGTAAAATCACAGACTACCGTAAGATGGCTGAAGGAGCTATGGAGCGCGTGGTTGACATCCTCAAAGCAGAATTTGAC
CGTAGCTTTAAATTGATCAATTCTAAAACTTACCCTGTTTCAGGTGGAGAATTGAACCCAGCAAATGTGGATTCAGAAATCG
AAGCCTTTGCGCAACTTGGAGTATCACAGTGGTTTGGATAGCAAGGAAGCTCACTATCTGGCAAATCTTTACGGTTCAAATGC
ACCGAAAGTCTTTGCACTTGCTCACAGCTTGGAACAACAGCACGCAAGACTCACTTTTATGCCTTCACTTATGCAA
GTATCGTTGAGCCAATTTTAGCCCAGTTGACTTCCTTCTTCGTCGTACCAATCACATGCTCTTTTATGCGTGATAGCTTGGATA
GTATCGTTGAGCCCAATTTTTGGATGAAATGGAACGAACTTACCGTGCTGATGT
CGAAGCAGCTCTCGCTAACAACGATTTTAGCAGAAAAAATTAAAAAATTAA

4106.8 (SEQ. ID. NO. 216)

4106.10 (SEQ. ID. NO. 217)

4107.1 (SEQ. ID. NO. 218)

4107.2 (SEQ. ID. NO. 219)

TGTTTATGGCGCTATTAAGCAGGATCCTCAATTGATCCAGCAGTTAATTGAGCTTTACCATGAGATGACCAAATCTCAGATGA GTTTTTTGGACTTGGAGAATTTAACAGATGAGGATAAGAGGGCGGATTTACTCTTGATTTTTGAGAAAGTAACAGCCTATCTT GTTGAGATTGTTATCGGGGCTTATGCTAGTAAGAAAGCCTATACCAGTCCTTTTAGCGAGGGCAATCTCTACCAAGCCAGCGT AAAATTTCTCCATCATCTGGCTTCTAAATACCAAACGCCTGCTCAGGACTGTTCTCAAACTCATGAGAAGATGGATAGTTTTG ACAAGGCCTCTCGTTTGTTGGAGTCTTCTTATGACTTTTCAGAACTCGCTTTGGATGTCGATGAGAAAGACCGTGAAAATTTA CAAATCTGGTCTTGTTTGACGCAAAAGGAGGAGTTGGAGCTAGTAGCCCGTAGTATTCGTCAGAAATTACATGAGAACTCAG ACCTGAGCTACAAGCATTTTCGTATTCTCTTGGGGGATGTAGCTTCTTACCAGTTATCTCTCAAAACCATTTTTGACCAGTATC AGATTCCTTTTTATCTTGGTAGAAGCGAAGCCATGGCTCATCATCCCTTGACTCAGTTTGTCGAGTCTATTTTAGCTTTAAAAC TTGAGCAATATATCCGCTATCTTGGTATCAATGGCTTGCCAGCCTTTCAGCAAACCTTCACCAAATCCCACCATGGAAAATTT AATCTTGAGCGTTTGAATGTCCTCCGCCTGAGAATTTTAGCACCTCTTGAAACCCTCTTTGCCAGCCGAAAACAAAAGGCTGA AAAACTCCTACAAAAATGGAGTGTCTTTCTAAAAGAAGGAGCTGTGACCAAGCAGTTACAAGATTTGACAACCACTTTGGAA GCTGTAGAACAGGAAAGACAAGCCGAAGTTTGGAAGGCTTTCTGCCATGTTTTAGAACAATTTGCGACTGTTTTTGCTGGTTC ACAGGTTAGTCTGGAAGACTTCCTAGCCTTGCTCCATTCTGGAATGAGTTTGTCCCAATACCGTACCATTCCAGCAACAGTGG ACACTGTTCTGGTGCAGAGTTACGATTTGATTGCACCATTGACTGCTGACTTTGTCTATGGTATTGGACTAACTCAGGACAAT TTACCAAAAATTTCTCAAAACACCAGTCTTCTGACAGATGAAGAAAGGCAAAACCTAAACCAAGCGACCGAAGAAGGCGTT CAATTACTGATTGCCAGCAGTGAAAATCTCAAGAAAAATCGCTACACTATGCTTTCCTTGGTCAATTCTGCTCGTAAGCAGTT GTTCTTGTCGGCTCCAAGCCTTTTTAACGAAAGTGAAAGTAAGGAATCTGCCTATCTTCAAGAGTTGATCCATTTTGGATTTA GGCGGAGAGAGAGAGGATGAATCACAAAGGACTGTCTAAGGAGGATATGGGGTCCTATCACAGTCTTTTGTCTAGTCTGGT TGCCTATCACCAGCAGGGTGAGATGAGCGATACTGAGCAAGATTTGACTTTTGTCAAGGTTCTGTCGCGTGTCATAGGTAAA AAACTAGATCAGCAAGGTCTGGAAAATCCAGCTATCCCAACCAGTCCAAGCAGCAAGACCTTAGCCAAGGACACCTTGCAA

4107.3 (SEQ. ID. NO. 220)

ATGAAGCTTATTCCCTTTTTAAGTGAGGAGGAGATTCAAAAACTGCAAGAAGCAGAAGCAAATTCGAGCAAGGAACAGAAG
AAAACTGCCGAGCAAATCGAAGCTATCTACACTTCTGCCCAGAATATCCTGGTCTCAGCATCGGCTGGTTCTGGAAAGACCT
TTGTCATGGCAGAGCGCATTCTGGACCAATTGGCGCGTGGTGTCGAAATTTCTCAACCTTTATCTCAACCTTTACCGTCAAG
CCTGCAACTGAACTTAAAGACCGTTTAGAGAAAAAAAACCAAGAAAAATCCAAGAAACAGATGATGTCGACCTCAAACAA
CACTTGGGTCGCCAGTTGGCAGACCTACCCAACGCTGCCATTGGAACCATGGATTCTTTCACACAAAAATTCCTTGGCAAAC
ATGGTTATCTGCTTGATATTGCACCTAATTTCCGTATTTTACAAAACCAAAGCGAGCAACTTATTCTCGAAAACGAAGTCTTT
CATGAGGTCTTTGAAGCGCATTACCAAGGTAAACAGAAGAGACCTTTAGTCATTTTGCTGAAAAACTTTGCTGGCAGAC
AGGACGAACGGGGTCTGCGCCAGCAGGTCTATAAAATCTATGACTTCCTCCAATCCACCAGTAATCCTCAAAAGTGGCTGAG
TGAATCTTTCCTCAAAGGATTTTACCAGTGATATTTACCAGTGAAAAAACGACCCTT
TGGGATTTTGGAAAACTTTTCCGTTACCATCTGGATAACGATGCCAAGGAGTTTGCAAAGCCTCATTTAGAAAATGTTCA
GGTTAATTCTGGATGAAATTGGCTCCCTAAATCAGGAGTCCGATAGCCAGGCTTTTCCAAGGCTGCCTATTTAGAAAATGTTCA
TCTCTAAAGGAAAAAACGGTCGAGCTCTGACTAAATCCAGCCGTAAGCCTGATTTTGAAACCACGA
AGAGAAAAAACGGTCGAGCTCTGACTAATGCCAGCCGTAAGGCTGATTTGAAGCCCCTGGCTGATGCCCAACGA
AGAGAAAAAGACCCCTTTTGCTAAACTAGGACCAGCCGTAAGGCTGATTTTGAAGCCCCTGGCTGATGCCCAACGA
AGAGAAAAAAGACCCCTTTTGCTAAACTAGGACCAATTATCAGACCAGCATTTTGAAGCCCCTGGCTGATGCCTACAACGA
AGAGAGAAAAAGACCCCTTTTTTTCCTTAAACTAGGACAATTATCAGACCAGCCGTAATTTGAAACCAACGA

4109.1 (SEQ. ID. NO. 221)

4110.2 (SEQ. ID. NO, 222)

4112.2 (SEQ. ID. NO. 223)

ATGAGTACTTTAGCAAAAATAGAAGCGCTCTTGTTTGTAGCGGGTGAAGATGGGATTCGGGTCCGCCAGTTAGCTGAACTCC
TCTCTCTGCCACCGACAGGCATCCAGCAAAGTTTAGGAAAATTAGCCCAGAAGTATGAAAAGGACCCAGATTCCAGTTTGGC
TTTGATTGAGACAAGTGTGCTTATAGATTGGTGACCAAGCCTCAATTTGCAGAGATTTTGAAGGAATACTCTAAGGCGCCT
ATCAACCAGAGCTTTGTCTCGGGCTGCCCTTGAGACCTTGTCCATTATTGCCTACAAACAGCCGATTACGCGGATAGAAATTGA
TGCCATCCGTGGAGTTAACTCGAGTGGAGCCTTGGCAAAGTTGCAGGCTTTTGACCTGATAAAGGAAGACGGGAAAAAGGA
AGTATTGGGGCGCCCCAACCTCTATGTGACTACGGATTATTTCCTAGATTACATGGGGATAAACCATTTAGAAGAAATTACCA
GTGATTGATGAGCTTGAGATTCAAGCCCAAGAAAGCCAATTATTTGGTGAAAGGATAGAAGAAGATGACAATAA

4113.1 (SEQ. ID. NO. 224)

4117.1 (SEQ. ID. NO. 225)

4119.2 (SEQ. ID. NO. 226)

4119.3 (SEQ. ID. NO. 227)

4119.4 (SEQ. ID. NO. 228)

4120.1 (SEQ. ID. NO. 229)

ATGTCAAAGATGGATGTTCAGAAAATCATTGCACCGATGATGAAGTTTGTGAATATGCGTGGCATTATAGCTCTAAAAGATG GGATGTTAGCAATTTTGCCATTGACAGTAGTTGGTAGTTTGTTCTTGATTATGGGACAATTGCCGTTCGAAGGATTAAATAAG AGCATTGCTAGTGTTTTTGGAGCTAATTGGACAGAGCCGTTTATGCAAGTATATTCAGGAACTTTTTGCTATTATGGGTCTAAT TTCTTGTTTTTCAATTGCCTATTCTTATGCTAAGAATAGCGGAGTAGAGGCTTTACCAGCTGGAGTTCTATCTGTATCTGCATT CTTTATTTTGCTAAGATCATCTTATATCCCTAAACAAGGTGAGGCGATTGGGGACGCTATTAGTAAAGTTTGGTTTGGAGGCC AAGGAATTATCGGTGCTATCATTATAGGTTTGGTAGTAGGAAGTATTTATACCTTCTTTATAAAGAGAAAAATTGTTATTAAG ATGCCAGAACAAGTTCCACAAGCTATTGCCAAACAGTTTGAAGCAATGATTCCAGCATTTGTAATTTTCTTATCTTCTATGAT TGTATATTTTTAGCGAAGTCATTGACTAATGGCGGAACATTCATAGAAATGATTTATTCTGCTATTCAAGTTCCGTTGCAAG GTTTAACTGGATCTTTGTATGGTGCTATTGGAATTGCATTCTTTATATCATTTTTGTGGTGGTTTGGTGTTCATGGGCAATCGG TAGTAAATGGAGTAGTGACAGCTCTGCTTTTATCTAATCTTGATGCTAATAAAGCTATGTTAGCCTCTGCTAATCTATCATTA GAAAATGGTGCACATATTGTTACTCAACAATTTTTAGATTCATTTTTAATTCTATCAGGTTCAGGGATTACGTTTGGTCTTGTA GTTGCCATGCTTTTTGCAGCAAAATCAAAACAATACCAAGCCTTAGGAAAAGTTGCAGCTTTTCCAGCAATATTTAACGTAA ATGAGCCAGTTGTATTTGGATTTCCGATTGTCATGAATCCAGTTATGTTTGTACCTTTCATTCTTGTTCCTGTACTTGCAGCTGT GATAGTATATGGAGCTATTGCAACAGGTTTCATGCAGCCATTCTCAGGGGTAACATTGCCTTGGAGTACACCAGCTATTTTAT CAGGATTTTTGGTGGGTGGATGGCAAGGAGTTATTACTCAGCTGGTGATATTAGCGATGTCTACATTGGTTTATTTTCCATTCT TTAAAGTACAGGATCGTTTAGCTTACCAAAATGAAATCAAACAATCTTAG

4121.2 (SEQ. ID. NO. 230)

ATGAAGAAAAAGGACTTAGTAGACCAACTAGTCTCAGAGATCGAGACGGGGAAAGTCAGGACACTGGGAATATACGGTCATGGAGCTCAGGGACACTAGAGACACTTTGCACAGGAATTGTACCAAGCTTTAGATTCTACTACAGTAAATTTGCTAGAGACAGATCCTTATATCACCTCAGGACGCCATCTGGTAGTACCCAAGGACGCCGCAATCAAAAGGTGACAGCCAGTCTGCCAGTGGCGCATGAACTGGAGAGTTTGCAGAGAGATATCCTTgCTTGCAGGCGGGTATGGATGTCTTGA

4122.1 (SEQ. ID. NO. 231)

4125.6 (SEQ. ID. NO. 232)

CTCATAAAATTGGAGATGCGGATGAATTTAAGCATGATACGGGTGTTGTCTATGCAGATTCTCCATTTATTCTTTTCTATTTTCACCTAAGAATTCTGATTATGATACGATTTCTAAGATAGCCAAGGATGTTTATGAGGTTCTAAAATGA

4125.7 (SEQ. ID. NO. 233)

4125.10 (SEQ. ID. NO. 234)

4126.1 (SEQ. ID. NO. 235)

ATGAAGCGTTCTTCTCTTTTAGTTAGAATGGTTATTTCCATCTTTCTGGTCTTTCTCATTCTCCTAGCTCTGGTTGGAACTTTCT ACTATCAATCAAGTTCTTCAGCCATTGAGGCCACCATTGAGGGCAACAGCCAAACGACCATCAGCCAGACTAGCCACTTTAT TCAGTCTTATATCAAAAAACTAGAAACCACCTCGACTGGTTTGACCCAGCAGACGGATGTTCTGGCCTATGCTGAGAATCCC AGTCAAGACAAGGTCGAGGGAATCCGAGATTTGTTTTTGACCATCTTGAAGTCAGATAAGGACTTGAAAACTGTTGTGCTGG TGACCAAATCTGGTCAGGTCATTTCTACAGATGACAGTGTGCAGATGAAAACTTCCTCTGATATGATGGCTGAGGATTGGTA CCAAAAGGCCATTCATCAGGGAGCTATGCCTGTTTTGACTCCAGCTCGTAAATCAGATAGTCAGTGGGTCATTTCTGTCACTC AAGAACTTGTTGATGCAAAGGGAGCCAATCTTGGTGTGCTTTCGTTTGGATATTTCTTATGAAACTCTGGAAGCCTATCTCAAT CAACTCCAGTTGGGGCAGCAGGGCTTTGCCTTCATTATCAATGAAAACCATGAATTTGTCTACCATCCTCAACACAGGTTTA TAGTTCGTCTAGCAAAATGGAGGCTATGAAACCCTACATCGATACAGGTCAGGGTTATACTCCTGGTCACAAATCCTACGTC AGTCAAGAGAAGATTGCAGGAACTGATTGGACGGTGCTTGGCGTGTCATCATTGGAAAAGTTAGACCAGGTTCGGAGTCAGC CTCCTTTGAAGGATTTGAGAGAAACCATGTTGGAAATTGCTTCTGGTGCTCAAAATCTTCGTGCCAAGGAAGTTGGTGCCTAT GAACTGAGAAGTAACTCGCCAATTTAATGCTATGTTGGATCAGATTGATCAGTTGATGGTAGCTATTCGTAGCCAGGAAG AAACGACCCGTCAGTACCAACTTCAAGCCCTTTCGAGCCAGATTAATCCACATTTCCTCTATAACACTTTGGACACCATCATC TGGATGGCTGAATTTCATGATAGTCAGCGAGTGGTGCAGGTGACCAAGTCCTTGGCAACCTATTTCCGCTTGGCGCTCAATCA AGGCAAGGACTTGATTTGTCTCTCTGACGAAATCAATCATGTCCGCCAGTATCTCTTTATCCAGAAACAACGCTATGGAGATA AGCTGGAATACGAAATTAATGAAAATGTTGCCTTTGATAATTTAGTCTTACCCAAGCTGGTCCTACAACCCCTTGTAGAAAAT GCTCTTTACCATGGCATTAAGGAAAAGGAAGGTCAGGGCCATATTAAACTTTCTGTCCAGAAACAGGATTCGGGATTGGTCA TCCGTATTGAGGATGATGGCGTTGGCTTCCAAGATGCTGGTGATAGTAGTCAAAGTCAACTCAAACGTGGGGGAGTTGGTCT AAAGTTGAAATATATATAAATAGAATAGAAACTAGCTAA

4126.7 (SEO. ID. NO. 236)

ATGAAGCGTTCTTCTCTTTTAGTTAGAATGGTTATTTCCATCTTTCTGGTCTTTCTCATTCTCCTAGCTCTGGTTGGAACTTTCT ACTATCAAGTCAAGTTCTTCAGCCATTGAGGCCACCATTGAGGGCAACAGCCAAACGACCATCAGCCAGACTAGCCACTTTAT TCAGTCTTATATCAAAAAACTAGAAACCACCTCGACTGGTTTGACCCAGCAGACGGATGTTCTGGCCTATGCTGAGAATCCC AGTCAAGACAAGGTCGAGGGAATCCGAGATTTGTTTTTGACCATCTTGAAGTCAGATAAGGACTTGAAAACTGTTGTGCTGG TGACCAAATCTGGTCAGGTCATTTCTACAGATGACAGTGTGCAGATGAAAACTTCCTCTGATATGATGGCTGAGGATTGGTA CCAAAAGGCCATTCATCAGGGAGCTATGCCTGTTTTGACTCCAGCTCGTAAATCAGATAGTCAGTGGGTCATTTCTGTCACTC AAGAACTTGTTGATGCAAAGGGAGCCAATCTTGGTGTGCTTCGTTTGGATATTTCTTATGAAACTCTGGAAGCCTATCTCAAT CAACTCCAGTTGGGGCAGCAGGGCTTTGCCTTCATTATCAATGAAAACCATGAATTTGTCTACCATCCTCAACACACAGTTTA TAGTTCGTCTAGCAAAATGGAGGCTATGAAACCCTACATCGATACAGGTCAGGGTTATACTCCTGGTCACAAATCCTACGTC AGTCAAGAGAAGATTGCAGGAACTGATTGGACGGTGCTTGGCGTGTCATCATTGGAAAAGTTAGACCAGGTTCGGAGTCAGC CTCCTTTGAAGGATTTGAGAGAAACCATGTTGGAAATTGCTTCTGGTGCTCAAAATCTTCGTGCCAAGGAAGTTGGTGCCTAT GAACTGAGAGAAGTAACTCGCCAATTTAATGCTATGTTGGATCAGATTGATCAGTTGATGGTAGCTATTCGTAGCCAGGAAG AAACGACCCGTCAGTACCAACTTCAAGCCCTTTCGAGCCAGATTAATCCACATTTCCTCTATAACACTTTGGACACCATCATC TGGATGGCTGAATTTCATGATAGTCAGCGAGTGGTGCAGGTGACCAAGTCCTTGGCAACCTATTTCCGCTTGGCGCTCAATCA AGGCAAGGACTTGATTTGTCTCTCTGACGAAATCAATCATGTCCGCCAGTATCTCTTTATCCAGAAACAACGCTATGGAGATA AGCTGGAATACGAAATTAATGAAAATGTTGCCTTTGATAATTTAGTCTTACCCAAGCTGGTCCTACAACCCCTTGTAGAAAAT GCTCTTTACCATGGCATTAAGGAAAAGGAAGGTCAGGGCCATATTAAACTTTCTGTCCAGAAACAGGATTCGGGATTGGTCA TCCGTATTGAGGATGATGGCGTTGGCTTCCAAGATGCTGGTGATAGTAGTCAAAGTCAAACTCAAACGTGGGGGAGTTGGTCT AAAGTTGAAATATATAAATAGAATAGAAACTAGCTAA

4127.4 (SEQ. ID. NO. 237)

4127.5 (SEQ. ID. NO. 238)

ATGGATATTAGACAAGTTACTGAAACCATCGCCATGATTGAGGAGCAAAACTTCGATATTAGAACCATTACCATGGGGATTTCTTTTTGGACTGTATCGATCCAGATATCAATCGTGCTGCTGCGGAGAAAATCTATCAAAAAAATTACGACAAAGGCGGCTAATTTAGTAGCTGTTGGTGATGAAATTGCGGCTGAGTTGGGAATTCCTATCGTTAATAAGCGTGTATCGGTGACACCTATTTCTCTGATTGGGGCAGCGACAGATGCGACGGACTACGTGGTTCTGGCAAAAGCGCTTGATAAGGCTGCGAAAGAGATTGGTGTGTGGACTTTATTGGTGGTTTTTCTGCCTTAGTACAAAAAGGTTATCAAAAAGGGAGATTCCATCAATTCCATTCCTCGCGCTTTG

4128.1 (SEQ. ID. NO. 239)

4128.2 (SEQ. ID. NO. 240)

ATGTCTAAATTTAATCGTATTCATTTGGTGGTACTGGATTCTGTAGGAATCGGTGCAGCACCAGATGCTAATAACTTTGTCAA
TGCAGGGGTTCCAGATGGAGCTTCTGACACACTGGGACACATTTCAAAAACAGTTGGTTTGAATGCCCAAACATGGCTAAA
ATAGGTCTTGGAAATATTCCTCGTGAAACTCCTCTTTAAGACTGTAGCAGCTGAAAGCAATCCAACTGGATATGCAACAAAAT
TAGAGGAAGTATCTCTTGGTAAGGATACTATGACTGGACACTGGGAAATCATGGGACTCAACATTACTGAGCCTTTCGATAC
TTTCTGGAACGGATTCCCAGAAGAAATCCTGACAAAAATCGAAGAATTCTCAGGACGCAAGGTTATTCGTGAAGCCAACAA
CCTTATTCAGGAACGGCTGTTATCTATGATTTTGGACCACGTCAGATGGAAACTGGAGAGTTGATTATCTTATCTTCAGCTGA
CCCTGTTTTTGCAGATTGCTGCCCACGAAGACATTATTCCTTTGGATGAATCACTGTGAATCTCTGTGAATTACC
CCTTGAGCGTCCTGCCCTTCTTGGTCGCATCATTTTCCTTTGGATGAACCAGGTAACTTCACTCGTACGGCAAACC
GTCGTGACTTGGCTGTATCTCCATTTTTCCCAACTGTTTTTGGATAAATTGAATGAGGCTGGTATCGATACCTTATGCTGTGGGTA
AAATCAACGATATCTTTAACGGTGCTGGTATCAACCATGACATGGGTCACAACAAGTCAAATAGTCATGGAATTGATACACT
ATTGAAGACTATGGGACTTGCTGAGTTTGAAAAAAGGATTCTCATTCACAAACCTAGTTGACTTTGATGCCCTTTTACGGCCATC
GTCGTAATGCTCACGGTTACCGTGAATTGCTTGCATGAGTTTGATGACGCTTACCTGAAATTATCCCTTTTTACGGCCATC
GTCGTAATGCTCACGGTTACCGTGAATTGCTTGCATGAGTTTGATGAACGCTTACCTGAAATTTTACCAGCACTATTGCTGGCGATACTTTTTACGGCCATC
GTCGTAATGCTCACGGTTACCGTGAATTGCATGAGTTTGATGAACGCTTACCTGAAATTATCCACTTGGTGGCCATC
GTCGTAATGCTCACGGGACATGGAAATGACCCAACGTATGCAGGAACGGATCACACTCGGGAATATATTCCATTGTT
GGCCTATAGCCCTGCCTTTAACGGGAAATGACCCAACGTATGCAGGAACAACTTTTCCAGCACCTGTTGCCGGATAACT
TTGGTGTGGAAACTGCTTATGATTGGGGAAAGGTTTCTTTAGGTATAAA

4129.2 (SEQ. ID. NO. 241)

ATGTTTATTTCCATCAGTGCTGGAATTGTGACATTTTTACTAACTTTAGTAGAAATTCCGGCCTTTATCCAATTTTATAGAAAG
GCGCAAATTACAGGCCAGCAGATGCATGAGGATGTCAAACAGCATCAGGCAAAAGCTGGGACTCCTACAATGGGAGGTTTG
GTTTTCTTGATTACTTCTTTTTGGTTGCTTTCTTTTTTCGCCCTATTTAGTAGCCAATTCAGCAATAATGTGGGAATGATTTTGT
TCATCTTGGTCTTGATTATGGCTTGGTCGGATTTTTTAGATGACTTTCTCAAGGGCTTTTCGTAAAATCAATGAGGGGCTTAATCCTA
AGCAAAAATTAGCTCTTCAGCTTCTAGGTGGAGTTATCTTCTATCTTTTCTATGAGCGCGGTGGCGATATCCTGTCTTTTG
GTTATCCAGTTCATTTTGGGATTTTTCTATATTTTTCTTTTCTGCCTAGTGGGTTTTTTCAAACGCAGTAAACTTGACAGA
CGGTGTTGACGGTTTAGCTAGTATTTCCGTTGTGATTAGTTTGCCTATGGAGTTATTGCCTATGTGCAAGGTCAGTAGA
ATATTCTTCTAGTGATTCTTGCCATGATTGGTGGTTTGCTCGGTTTCTTCATCTTTTAACCATAAGCCTGCCAAGGTCTTTATGG
GTGATGTGGGAAGTTTGGCCCTAGGTGGGATGCTGGCAGCTATCTCTATGGCTCCCCCACAAGAATGGACTCTCTTGATTATC
GGAATTGTTATGTTTTTTGGAACAACTTCTGTTATGATCAGCTCAGTTATTTCAACCTGACAGGTGGTAAACGTATTTTCCG
TATGACGCCTGTACATCACCATTTTGAGCTTGGGGGATTTGCTCTGGTAAAGGAAATCCTTGGAGCGAGTGGAAGGTTGACTTCT
TCTTTTGGGGAGTTGGACATCACCATTTTTAGCAAGTTGCTCTGGTAAAGGAAATCCTTTGAGCAGGTGGAAGGTTGACTTCT
TCTTTTGGGGAGTTGGAGAGTTGACAAGCTCCCTAGCAATTTTTATTTTATTTTGATTAA
TTTTTTGAGCAGCTGGAAGTTGACAAGCTCCCTGACCCTAGCAATTTTTATTTTATTTTGATTAA
TTTTTTAGATGACGCAGTTGACATCACCATTTTAACCAAGTTGACTTCATTATTTTATTTTTATTTTATTTTATATTTTAACTTAA

4133.1 (SEQ. ID. NO. 242)

TTGTTTAAGAAAAATAAAGACATTCTTAATATTGCATTGCCAGCTATGGGTGAAAACTTTTTGCAGATGCTAATGGGAATGGT GGACAGTTATTTGGTTGCTCATTTAGGATTGATAGCTATTTCAGGGGTTTCAGTAGCTGGTAATATTATCACCATTTATCAGG CGATTTTCATCGCTCTGGGAGCTGCTATTTCCAGTGTTATTTCAAAAAGCATAGGGCAGAAAGACCAGTCGAAGTTGGCCTAT CATGTGACTGAGGCGTTGAAGATTACCTTACTATTAAGTTTCCTTTTTAGGATTTTTTGTCCATCTTCGCTGGGAAAGAGATGAT GGTTTAATGACTAGTCTAGGAGCCTTGATTCGTGCAACGCATAATCCACGTCTGCCTCTCTATGTTAGTTTTTTATCCAATGCC TTGAATATTCTTTTTTCAAGTCTAGCTATTTTTGTTCTGGATATGGGGATAGCTGGTGTTGCTTGGGGGGACAATTGTGTCTCGT TTGGTTGGTCTTGTGATTTTGTGGTCACAATTAAAACTGCCTTATGGGAAGCCAACTTTTGGTTTAGATAAGGAACTGTTGAC CTGTTCCTCATGTTGCCCCTGTCCTTTAGTATATGTCTTGGGTGTACCATTAACTCATCTCTATACGACTGATTCTCTAGCG GTGGAGGCTAGTGTTCTAGTGACACTGTTTTCACTACTTGGGACCCCTATGACGACAGGAACAGTCATCTATACGGCAGTCTG ATGGGGATTGTGCTTGGTTGGGGCTTGCCTGGTATTTGGGCAGGGTCTCTCTTGGATAATGGTTTTCGCTGGTTATTTCTACGC TATCGTTACCAGCGCTATATGAGCTTGAAAGGATAG

4135.2 (SEQ. ID. NO. 243)

ATGCAAACTCAAGAAAAACACTCGCAAGCAGCCGTTCTTGGCTTGCAGCACTTACTAGCCATGTACTCAGGATCTATCCTGG GTGGCAACCTTCCTCCAACTCCAACTCAACAAATACTTTGGGATTGGACTCCCAGTCGTTCTTGGAGTTGCATTCCAGTCGGT CGCTCCCTTGATTATGATTGGGCAAAGCCATGGTAGTGGCGCTATGTTTGGTGCCCTTATCGCATCTGGGATTTACGTGGTTC TTGTTTCAGGCATCTTCTCAAAAGTAGCCAATCTCTTCCCATCTATCGTAACAGGATCTGTTATTACCACGATTGGTTTAACCT CTGATTATCCTCTTGATCAACATCTTTACCAAAGGATTTATCAAGTCTATCTCTATTTTGATTGGTCTGGTTGTTGGAACTGCC ATTGCTGCTACTATGGGCTTGGTGGACTTCTCTCTGTTGCGGTAGCTCCACTTGTCCATGTCCCAACTCCACTCTACTTTGGG ATGCCAACCTTTGAAATCTCATCTATTGTCATGATGTGTATCATCGCAACGGTGTCTATGGTTGAGTCAACTGGTGTTTATCTG GCCTTGTCTGATATCACAAAGGATCCAATCGACAGCACGCGCCTTCGCAACGGATACCGCGCAGAAGGTTTGGCCGTACTTC TCGGAGGAATCTTTAACACCTTCCCTTACACCGGATTTTCACAAAACGTTGGTTTAGTTAAATTGTCAGGCATCAAAAAACGC CTGCCAATCTACGCAGCTGGTTTCCTGGTTCTCCTTGGACTGCTTCCTAAGTTTTGGCGCCCCTTGCCCAAATCATTCCAAGC TCCGTCCTCGGTGGTGCCATGCTGGTAATGTTTGGTTTTGTATCAATTCAAGGGATGCAAATCCTCGCCCGTGTTGACTTTGCT AACAATGAACACAACTTCCTTATCGCAGCTGTTTCAATCGCTGCAGGTGTCGGTCTCAACAACAGTAATCTCTTTGTCAGCAT ATAAAAAGAAATAA

4136.2 (SEQ. ID. NO. 244)

4137.2 (SEQ. ID. NO. 245)

TCTAAAAAAGAGATGAAAGGAAAAGAGTTCTTTAAAGGAGATGGTTCGGTTCCTTGGTATGTTACTTCGGTATCCATTTTTGC CACAATGCTCAGTCCGATTTCCTTCTTGGGACTCGCTGGTAGCTCTTATGCAGGTAGCTGGATTTTATGGTTTGCTCAATTAGG GATGGTAGTAGCTATTCCACTGACAATTCGTTTTATCTTACCTATCTTTGCACGGATAGACATCGATACGGCATATGATTACTT TATGTACCTCCCATCAGCTGGTTTATCAGTATTGACAGGAATTGACATCAATATTTTGATTATTTTGATGGGTGTAGTTGCAAT TGTTTATTCTTATACTGGTGGTCTAAAATCCGTATTATGGACAGACTTTATTCAAGGTGTGATTCTGATTAGTGGTGTCGTTTT AGCTTTATTTGTACTGATTGCTAATATTAAAGGTGGCTTTGGTGCAGTAGCAGAACATTAGCAAACGGGAAATTCCTTGCTG CAAATGAAAAACTTTTCGATCCTAACTTGCTTTCAAACTCCATCTTTTTAATTGTGATGGGTTCAGGCTTTACAATCTTGTCTT CCTATGCTTCATCTCAAGATTTGGTTCAACGTTTTACTACAACACAAAATATTAAGAAACTTAATAAGATGTTGTTCACAAAC GGTGTTTTGTCACTTGCAACTGCAACAGTCTTTTACTTGATTGGTACAGGCTTGTACGTATTCTATCAAGTACAAAATGCAGA TAGTGCAGCTAGCAATATCCCTCAAGACCAAATCTTTATGTACTTTATTGCATACCAGTTACCAGTAGGTATCACAGGTTTGA TCTTGGCAGCGATTTATGCAGCATCTCAATCAACTATTTCAACAGGTTTGAACTCTGTTGCAACTTCATGGACATTGGATATT CAAGATGTCATTTCTAAAAATATGTCAGACAATCGTCGTACGAAAATTGCACAATTCGTATCTCTAGCAGTAGGTTTATTCTC AATTGGTGTTTCCATTGTCATGGCTCACTCAGATATTAAATCTGCATACGAATGGTTCAATAGTTTCATGGGACTTGTACTTG GTCTACTTGGTGGTGTATTTATTCTTGGATTTGTTTCTAAAAAAGCAAATAAACAAGGTGCTTATGCAGCGCTGATTGTATCA ACCATCGTCATGGTATTTATTAAATACTTCCTTCCTACAACAGCTGTTAGCTACTGGGCATATTCATTGATTTCAATCTCTGTA TCAGTAGTTTCAGGTTATATTGTATCTGTTCTTACTGGAAATAAAGTATCTGCACCTAAATATACAACGATTCATGATATTAC AGAAATTAAAGCGGATTCAAGTTGGGAAGTTCGTCACTAA

4138.1 (SEQ. ID. NO. 246)

ATGAAATTTAGTAAAAAATATATAGCAGCTGGATCAGCTGTTATCGTATCCTTGAGTCTATGTGCCTATGCACTAAACCAGCA TCGTTCGCAGGAAAAATAATCGTGTCTCTTATGTGGATGGCAGCCAGTCAAGTCAGAAAAGTGAAAACTTGACA CCAGACCAGGTTAGCCAGAAAAGAAGGAATTCAGGCTGAGCAAATTGTAATCAAAATTACAGATCAGGGCTATGTAACGTCA

4139.1 <u>(SEQ. ID. NO. 247)</u>

4139.5 (SEQ. ID. NO. 248)

4139.8 (SEQ. ID. NO. 249)

4141.1 (SEQ. ID. NO. 250)

ATGATGAAGTTCATATTGGATATTGTTAGTACACCAGCTATTTTAGTAGCTTTAATTGCAATCTTAGGATTAGTTCTTCAGAA GAAGAAATTACCTGATATTAATAAAGGTGGAATTAAGACCTTTGTTGGTTTCTTAGTTGTATCTGGTGGTGCAGGAATTGTAC AAAATTCTTTAAATCCATTTGGTACCATGTTTGAGCATGCTTTTCATTTATCTGGCGTTGTGCCGAATAATGAAGCAATTGTAG CTGTAGCTTTAACAACATATGGCTCAGCTACTGCAATGATTATGTTTGCAGGCATGGTGTTCAATATCTTAATCGCTCGTTTTTA CTCGATTTAAATATATTTTTTTAACAGGGCACCACACTCTATATATGGCATGTATGATTGCGGTCATTTTATCAGTTGCTGGCT TTACTAGCTTGCCTCTCATCTTACTAGGAGGATTAGCACTCGGTATTATTATGAGTATTTCCCCAGCATTTGTGCAAAAATATA TGGTTCAATTAACTGGAAATGACAAGGTAGCTTTAGGTCATTTCAGTTCTTTGGGATATTGGTTGAGTGGTTTTACTGGTAGC CTTATCGGTGACAAATCAAAATCAACAGAGGACATTAAATTTCCAAAGAGTTTAGCTTTTTTACGTGATAGTACTGTTAGTAT TACTTTATCCATGGCAGTTATTTACATTATTGTAGCTATCTTTGCAGGGTCAGAATATATAGAAAAAGAAATCAGTAGTGGTA CAAGTGGTCTAGTTTATGCTTTACAATTAGCAGGTCAATTTGCAGCAGGGGTATTTGTTATTTTAGCAGGTGTTCGCCTTATTT TGGGCGAAATTGTTCCAGCCTTTAAAGGTATTTCAGAGCGTCTTGTACCTAATTCAAAACCTGCTTTGGATTGTCCGATTGTTT CAGGAACGGTTGTTATCTTACCAGGTGTTGTGCCTCATTTCTTCTGTGGAGCGACTGCAGGTGTCATTGGGAATGCATCTGGT GGTGTTCGTGGAGCCACTATTGGAGCATTTTTACAAGGTATTTTAATCAGTTTTCTTCCAGTCTTTTTAATGCCAGTTTTTGGGA GGACTTGGTTTCCAAGGATCAACTTTCTCAGATGCAGATTTTTGGTCTATCAGGAATTATTTTAGGAATGTTAAATCAATTTTGG CTCACAAGCAGCATTGTGATTGGTCTTGTTCTTATTCTAGCAGTTATGTTTGGAGTATCCTTTATTAAAAAGCCATCTGCAAC **GGAGGAATAA**

4142.3 (SEQ. ID. NO. 251)

4142.4 (SEQ. ID. NO. 252)

ATGAATCCAATCCAAAGATCTTGGGCTTATGTCAGCAGAAAGCGACTGAGAAGTTTTATTTTATTTTCTGATTTTATTTGGTCTT ATTGGCCGGAATTTCAGCCTGTTTGACTCTGATGAAGTCCAACAAAACAGTAGAAAGCAATCTTTATAAATCACTCAATACA TCTTTTTCTATTAAGAAGATAGAGAATGGTCAGACATTCAAGTTGTCAGACCTAGCATCTGTAAGCAAGATTAAGGGGCTGG AAAATGTCTCTCTGAACTTGAGACGGTCGCAAAACTAAAAGACAAGGAAGCAGTGACTGGCGAGCAGAGCGTGGAGCGTG ATGATTTATCAGCTGCAGACAATAACTTGGTTAGCTTAACGGCTCTTGAGGATTCATCCAAGGATGTAACCTTTACCAGTTCG GCTTTCAATCTAAAAGAAGGGCGACACCTTCAAAAAGGGGATTCCAAGAAAATCCTTATCCACGAAGAATTGGCTAAGAAG AACGGTCTTTCGCTTCATGACAAGATTGGCTTGGATGCTGGTCAGTCTGAATCTGGAAAAGGACAAACAGTAGAGTTTGAGA TTATCGGCATCTTTTCTGGTAAAAAACAAGAGAAATTCACAGGCTTGTCTTCTGACTTCAGTGAAAATCAAGTCTTTACAGAC TATGAAAGTAGCCAAACCCTTTTGGGCAATAGTGAAGCTCAAGTCAGTGCAGCACGCTTCTATGTAGAAAATCCTAAGGAAA TGGACGGACTCATGAAGCAGGTAGAAAACTTGGCCTTGGAAAATCAAGGCTACCAAGTCGAAAAGGAAAACAAGGCTTTTG AACAAATCAAAGACTCAGTTGCAACTTTCCAAACCTTCCTGACCATCTTCCTTTATGGGATGTTGATAGCAGGAGCTGGAGCC TTAATTCTGGTTTTGTCTCTCTGGTTGAGAGACGGGTCTATGAAGTGGGGATTTTACTTGCACTTGGAAAAGGCAAGAGCTC GATCTTCCTACAATTCTGTTTAGAGGTAGTTTTGGTATCTCTTGGAGCTTTGCTTCCAGCATTTGTTGCAGGAAACGCAATCAC ACTAGCATCTTATCTTTTGCAGAATCCTATGTTTTTCTAGTTCTGCTTAGTTGCTTATCTGTAGCCCTTTGTTTCCTATTCTTATT TAGAAAATCACCGAAAGAAATTTTATCATCTATTAGTTAA

4142.5 (SEQ. ID. NO. 253)

4144.1 (SEQ. ID. NO. 254)

4144.2 (SEQ. ID. NO. 255)

4144.3 (SEQ. ID. NO. 256)

ATGACAGAAACCATTAAATTGATGAAGGCTCATACTTCAGTGCGCAGGTTTAAAGAGCAAGAAATTCCCCAAGTAGACTTAA
ATGAGATTTTGACAGCAGCCCAGATGGCATCATCTTGGAAGAAATTCCCAATCCTACTCTGTGATTGTGGTACGAAGTCAAGA
GAAGAAAGATGCCTTGTATGAATTGGTACCTCAAGAAGCCATTCGCCAGTCTGCTGTTTTTCCTTCTCTTTTGTCGGAGATTTGA
ACCGAGCAGAAAAGGGAGCCCGACTTCATACCGACACCCTCCAACCCCAAGGTGTGGAAGGTCTCTTGATTAGTTCGGTCGA
TGCAGCTCTTGCTGGACAAAACGCCTTGTTGGCAGCTGAAAGCTTGGGTGGTGGTTGATTATCGGTTTGGTTCGATACA
AGTCTGAAGAAGTGGCAGAGCTCTTTAACCTGACTACACCTATTCTTTTTGGGATGGCACTGGGTGTGCCAAATCAA
CATCATGATATGAAACCGAGACTCCTAGAGAATGTTTGTCTTTTGAGGAAGAATACCAAGAACAGTCAACTGAGGCAATCC
AAGCTTATGACCGTGTTCAGGCTGACTATGCTGGGGCGCGTGCGACCACAAGCTGGAGTCAGCGCCTAGCAGAACAGTTTTGG
TCAAGCTGAACCAAGCTCAACTAGAAAAAATCTTGAACAGAAGAAATTATTGTAG

4146.1 (SEQ. ID. NO. 257)

4146.2 (SEQ. ID. NO. 258)

4147.1 (SEQ. ID. NO. 259)

4147.2 (SEQ. ID. NO. 260)

4147.3 (SEQ. ID. NO. 261)

4149.1 (SEQ. ID. NO. 262)

GTCTCGAAATCGGTCGTTTCAAGACAGGAACCCCTCCACGTGTCAAGGCTTCTTCTATCAATTACGATGTGACAGAAATTCAGCACGAGAGACGAAGTGCCTAATCATTTCTCATACACTTCACGTGATGAGGATTATGTCAAGGACCAAGTACCATGCTGGTTGACCTATACCAATGGTACCAGTCATGAGATTATCCAAAACAACCTCCACCGTGCGCCTATGTTTACAGGTGTGGTCAAGGAGTGGGCCTCGTTACTGTCCGTCGATTGAAGACAAGATTGTGCGCTTTGCGGACAAGGAACGTCACCAACTCTTCCTTGAGCCAGAGGGCGCAATACTGAGGAAGTCTATGTGCAAGGACTTTCAACCAGTCTGCCTGAGGATGTCCAGCGTGACTTGGTGCATTCCATCAAAGGTTTGGAAAATGCAGAGATGATGCGGACAGGTTATGCTATTGA

4149.2 (SEQ. ID. NO. 263)

4149.3 (SEO. ID. NO. 264)

4152.2 (SEQ. ID. NO. 265)

4154.1 (SEQ. ID. NO. 266)

ATGACTACTTTTAAAGATGGATTTTTATGGGGTGGTGCTGTTGCTGCTCATCAACTTGAAGGTGGATGGCAAGAAGGTGGCA AGGGAATTAGTGTTGCTGATGTTATGACTGCTGGTCGTCATGGAGTAGCTCGTGAAATTACTTTGGGAGTTTTAGAGGGTAAA TATTATCCAAATCATGAGGCGATAGATTTTTATCACCGTTATAAAGAAGATATAGCACTTTTTGCTGAAATGGGATTCAAGTG CTTCCGTACCTCTATTGCATGGACACGTATCTTTCCAAAAGGTGATGAGGTTAGAGCCGAATGAAGAAGGATTACAGTTTTATG ATAATCTTTTTGATGAATGCTTAAAGAATGGTATTGAACCTGTCATCACTCTATCTCATTTTGAAATGCCTTATCACTTAGTGA AAGGTTAAATATTGGATGACTTTCAATGAAATCAATAATCAAGCGAATTATCAGGAAGATTTTGCACCATTTACTAACTCAG GTATTGTATATGAGGAAGGTGATAATAGAGAAGCAATTATGTATCAAGCAGCACATTACGAATTAGTTGCTTCTGCACGAGC TGTAAAAATTGGTCATGAGATTAATCCAGATTTTCAAATAGGTTGTATGATTGCGATGTGTCCAATTTATCCAGTTACTTGCA ATCCTAAGGATATCTTAATGGCAATGAAAGCTATGCAGAAGCGTTATTATTTTTGCTGATGTGCATGTTTTAGGTAAATATCCT GAGCATATTTCAAGTATTGGGAACGAAAAGGTATTTCAGTTGATTTTACTGCCCAGGATAAAGAAGATTTACTTGGTGGGA CTGTAGATTACATTGGTTTCAGTTACTATATGTCCTTTGCTATCGACTCTCATCGTGAAAATAATCCTTATTTTGATTATCTTG AAACAGAAGATTTAGTGAAAAATAATTATGTTAAGGCTTCTGAATGGGAGTGGCAAATTGATCCAGAAGGTTTGCGTTATGC GTTAAATTGGTTTACAGACCACTATCACTTACCACTCTTTATTGTTGAAAATGGTTTTTGGAGCTATAGATCAAGTTGCAGCAG ATGGTATGGTACATGATGATTATAGAATTGAATATCTAGGTGCCCATATTCGTGAAATGAAAAAGGCTGTAGTTGAAGATGG TGTTGATTTAATGGGTTATACTCCATGGGGATGTATTGATTTGGTTTCAGCTGGTACCGGTGAAATGCGGAAACGTTATGGCT TTATTTATGTAGATAAAGATGATAATGGGAAGGGAAGTTATAATCGTTCCCCGAAAAAAATCTTTTGGCTGGTATAAGGAAGT **TATTTCATCTAACGGTGAATCAGTAGAATAG**

4154.2 (SEQ. ID. NO. 267)

4155.1 (SEQ. ID. NO. 268)

ATGAAAAATTITATGTAAGTCCAATTITTCCTATTCTAGTAGGATTGATTGCGTTTGGAGTCTTATCCACTITCATTATTITT TATACAAGGAGTGATGTAGAACAGATACAGTATGTAAACCACCAAGCGGAAGAAGTTTGACAGCTCTATTGGAACAGATG CCTGTAGGTGTTATGAAATTGAATTTATCTTCTGGAGAGGTTGAGTGGTTTAATCCCTATGCTGAATTGATTTTTGACCAAGGA AGATGGTGATTTTGATTTAGAAGCTGTTCAAACGATTATCAAGGCTTCAGTAGGAAATCCGTCTACTTATGCCAAGCTTGGTG AGAAGCGTTATGCTGTTCATATGGATGCTTCTTCCGGTGTTTTGTATTTTGTAGATGTATCCAGGGAACAAGCCATAACAGAT GAATTGGTAACAAGTAGACCAGTGATTGGGATTGTCTCTGTGGATAATTATGATGATTTGGAGGATGAAACTTCTGAGTCAG ATATTAGTCAAATCAATAGTTTTGTAGCTAATTTTATATCAGAGTTTTCAGAAAAACACATGATGTTTTCTCGTCGGGTAAGT GAAGAGTCGAAACAGAGACAGTTGCCCTTGACCTTAAGTATGGGATTTTCTTATGGCGATGGAAATCATGATGAGATAGGGA AAGTTGCTTTGCTCAATTTGAACTTGGCTGAAGTACGTGGTGGCGACCAGGTGGTTGTTAAGGAAAACGACGAAACGAAAAA TCCAGTTTATTTTGGTGGTGGTCTGCTGCTTCAATCAAGCGTACACGGACTCGTACGCGCGCTATGATGACAGCTATTTCAG ATAAGATTCGGAGTGTAGATCAGGTTTTTGTAGTCGGTCACAAAAATTTAGACATGGATGCTTTGGGCTCTGCTGTAGGTATG CAGTTGTTCGCCAGCAATGTGATTGAAAATAGCTATGCTCTTTATGATGAAGAACAAATGTCTCCAGATATTGAACGAGCTGT TTCATTCATAGAAAAAGAAGGAGTTACGAAGTTGTTGTTGTTTAAGGATGCAATGGGGGATGGTGACCAATCGTTCTTTGTTG CCATAGAAGGGATCAGGATTTTCCAGATAATGCGGTTATTACTTATATCGAAAGTGGTGCAAGTAGTGCCAGTGAGTTGGTA ATTGCTATCCAGGAAATCGCTGCGACAGATTTTGAAGAATATCGTGAGGTCAATGAACTTATTTTACAGGGGCGTAAATTAG GTTCAGATGTACTAATAGCAGAGGCTAAGGACATGAAATGCTATGATACAGTTGTTATTAGTAAGGCAGCAGATGCCATGTT AGCCATGTCAGGTATTGAAGCGAGTTTTTGTTCTTGCGAAGAATACACAAGGATTTATCTCTATCTCAGCTCGAAGTCGTAGTA AACTGAATGTACAACGGATTATGGAAGAGTTAGGCGGTGGAGGCCACTTTAATTTGGCAGCAGCTCAAATTAAAGATGTAAC

4156.1 (SEQ. ID. NO. 269)

ATGAAAGAGAAAATATGTGGAAAGAATTGTTGAATCGTGCAGGCTGGATTTTGGTCTTTTTACTTGCCGTCCTTTTATATCA GGTTCCCCTAGTGGTTACCTCTATTTTGACTTTAAAAGAAGTAGCCCTGCTACAGTCAGGGCTGATAGTTGCTGGCCTTTCAA TTGTGGTTCTGGCTCTATTTATTATGGGAGCTCGTAAAACCAAGTTAGCTAGTTTTAATTTTTCTTTTTTTAGAGCTAAAGATT TGGCACGTTTGGGCTTGAGTTATCTAGTTATTGTCGGGTCAAATATACTTGGTTCCATTTTATTGCAACTGTCAAATGAGACG CCGATTTGTGAGGAAATCTTGTGTGGGGATTGTTCCTAAAAAGATTTTCCGAGGCAAGGAGAACTTGGGATTTGTAGTCG GTACGATTGTGTTTGCTTTATTGCATCAACCAAGTAATTTACCTTCTTTATTGATTTATGGAGGTATGTCGACAGTTCTATCTT CAGGCTGGATTTTGGTCTTTTTACTTGCCGTCCTTTTATATCAGGTTCCCCTAGTGGTTACCTCTATTTTGACTTTAAAAGAAG AAGTTAGCTAGTTTTAATTTTTCTTTTTTTTAGAGCTAAAGATTTGGCACGTTTGGGCTTGAGTTATCTAGTTATTGTCGGGTCA AATATACTTGGTTCCATTTTATTGCAACTGTCAAATGAGACGACAACAGCTAACCAGTCTCAGATTAATGATATGGTTCAAAA AAAGATTTTCCGAGGCAAGGAGAACTTGGGATTTGTAGTCGGTACGATTGTGTTTGCTTTATTGCATCAACCAAGTAATTTAC CTTCTTTATTGATTTATGGAGGTATGTCGACAGTTCTATCTTGGACAGCCTACAAGACCCAACGTTTGGAAATGTCGATCTTG CTTCACATGATTGTTAATGGGATTGCTTTCTGTTTGTTGGCTCTTGTGGTGATTATGAGTCGGACATTAGGAATTTCTGTTTAA

4156.4 (SEQ. ID. NO. 270)

4157.2 (SEQ. ID. NO. 271)

ATGAAAGACTTGTTTTTTAAAGAGAAAGCAGGCCTTTCGTAAGGAGTGTCTTGGTTATCTGCGCTATGTCTCAATGACCACTT TGTCTTGTTCCTGCTTGTTCGTTGGGCTTTCTAGCCTACCAGTACAGTCAACTCTTACAACATTTTCCTGAAAATCATTGGCC TATCCTTTTGTTTGTTGAGAAATTACGTCTGTTTTACTTTTACTTTTGGGAGAGGAACTGCCACCTATATGGAGGCTCCAGACAAGCT CTTTCTTTAGTTGGAGAAAGAGGAAATTAAGCTCCATCTCAAGCGTCAAACTGGCATTTCCCTAGTCTTTTTGGCTCTTTTGTACA GACCCTTTTCTTCTGCTGTTATTTGCCCTTTTTTTTATTTTTTATTTTTTCACAATGGGTTAGCAACTGGACTGGACTGGACTGTGCTTTTATTTC TCAAGAAAGCAAGCGTAAGCAAGTCTTGCTTCTTTTTTCGCTCTTTTACCCTCTTTTACCCAGGTCAAGGGAAATTTCAAACAGCGTTAACCCGTCCTTTTACCCTCTTTTACCCTCTTTTACCCTCTTTTACCCAGGTCAAAGCGAAATCTCTATCTCACCGCTTCTTATTCACCAGAAGCCTAAAATCTCTATCTCGCGTTCTTATT

4158.1 (SEO. ID. NO. 272)

4158.2 (SEQ. ID. NO. 273)

4158.3 (SEQ. ID. NO. 274)

ATGTCCTCTTTTCGGATCAAGAATTAGTAGCTAAAACAGTAGAGTTTCGTCAGCGTCTTTCCGAGGGAGAAAGTCTAGACG
ATATTTTGGTTGAAGCTTTTGCTGTGGTGCGTGAAGCAGATAAACAGTAGAGTTTTCGTCAGCGTTTTCCTTATGATGTTCAAGTCATG
GGAGCTATTGTCATGCACTATGGAAATGTTGCTGAGATGAATACGGGGGAAGGTAAGACCTTGACAGCTACCATGCCTGTCT
ATTTGAACGCTTTTTCAGGAAGAAGGAGTGATGGTTGTGACCTCCTAATTACGATATTTATCAAAGCGTGATGCCGAGGAAATGGG
TCAAGTTTATCGTTTTCTAGGATTGACCATTGGTTGTACCATTTACGGAAGATCCAAAGAAGGAGATGAAAGCTGAAGAAAAG
AAGCTTATCTTCTGGGATATCATCTACACAACCAATAGTAATTTAGGTTTTTGATTACTAAATGATAACCTAGCCTCGAA
TGAAGAAGGTAAGTTTTTACGACCGTTTAACTATGTGATTATTGATAGAAATTGATAATCTTTGATAAGTGCACAAAACTC
CTCTGATTATTGCGGGTTCTCCTCGTGTTCAGTCTAATTACTATGCGATCATTGATACACTTTGTAACAACCTTTGGTCGAAGA
GAGGATTATATCTTTAAAGAGGAGAAAGAGGAGGTTTGGCTCACTACTAAGGGGGCCCAAGTCTGCTGAGAATTTCCTAGGGA
TTGATAATTTATACAAGGAAGAAGAGAGGATGTTTTTGCTCGTCATTTTGGTTTTATTGCGATTCGAGCTCATAAGCTCTTTACTAAA
GATAAGGACTATATCATTCGTGGAAATGAGGATGGTACTGGTTGATAAAGGGAACAGGCGTCTAATGGAAATGACTAAACTTC
AAGGAGGTCTCCATCAGGCTATTGAAGCCAAGGAACATCTCAAAATTATCTCCTGAGAACGGGCTCTAATGGCCTCGATCACCTA
TCAGAGTCTCTTTTAAGAATGTTTAATAAGATATCTGGTATGACAAGGGACAGGGACAGGGCGTATTGGCCTCGATCACCTA
TCAGAGTCTTTTTTAAGAATGTTTAATAAGATATCTCGGTATGACAAGGACACGGACAAGGACACGGGAAAAAAGAGTTTAATTGAAACT
TCAGAGTCTTTTTTAAGAATGTTCAACCAATCGTCCGAGACAACGGA
TACAATATGTCTGTAGTACCGCATTCCAACCAATCGTCCGAGACAACGGA
TACAATATGTCTGTAGTACCACCAATCCGTCCGAGACAACGGA
TACAATATGTCTGTAGTACCACATTCCAACCAATCGTCCCAGACACGGA
TACAATATGTCTGTAGTACCACTTCCAACCAATCGTCCCAGAGCAACGGA
TTACAATATGTCTGTAGTACCACTTCCAACCAATCCTCCCAGACACACGGA
TACAATATGTCTGTAGTACCACTTCCAACCAATCCTCCCAGACACACCGA

4158.4 (SEQ. ID. NO. 275)

AGAATTGCTAAAAACTATCGCTGACGTGACTTCTAAAATTGTCGAATGGATCATCAATCTGGCTCCATTTGGAATCCTTGGTC TTGTTTTTAAAACCATTTCTGACAAGGGAGTCGGAAGCCTTGCCAACTACGGTATTTTATTGGTTCTATTAGTAACGACTATG CTTTTTGTTGCCCCTGTGGTCAACCCTTTGATTGCCTTCTTTATGAGACGCAATCCTTACCCTCTAGTTTGGAACTGCCTCC GTGTCAGCGGTGTGACAGCCTTTTTCACTCGTAGTTCTGCGACTAACATTCCTGTCAACATGAAACTCTGCCATGACCTTGGA CTCAACCCAGATACCTATTCTGTTTCTATCCCACTCGGTTCTACTATCAATATGGCTGGAGTAGCGATTACCATTAACCTTTTG ACCCTTGCTGCAGTTAACACTCTTGGAATTCCTGTTGACTTTGCCACAGCCTTTGTCCTCAGTGTGGTAGCAGCTATCTCATCC TGTGATGCTTCAGGTATTGCCGGAGGTTCCCTCCTTCTTATCCCAGTTGCTTGTAGCCTTTTCGGTATTTCTAACGATATTGCC ATACAAATTGTTGGGGTTGGTTTTGTGATTGGTGTCATCCAAGACTCATGTGAAACAGCCCTTAACTCTTCTACAGATGTCCT CTTTACCGCCGTTGCCGAATACGCAGCAACCCGTAAAAAATAA

4158.5 (SEQ. ID. NO. 276)

GCAGTTTCGGCTGGAATTATCGCTCTCTTGAGCCTATCTGATACGCGTAGAAGTACTTTAAAACTGGCTCGCAATCGTCTTTTT TCTATGCTTCTAGCTCTGGCTATCGGTGTTCTAGCTTTTCACTTGAGCGGATTTCATATCTGGAGTCTCGGCCTCTATCTGGCC TTCTACGTTCCTTTAGCCTACAAGATGGGCTGGGAAATTGGCATCACACCAAGCACTGTTTTGGTTAGCCATCTCTTTGGTTCA TATGCCTTCACGAGAAGAGGAAATCCAGCACTACCACACGCTGGTGGAAGAAAGTTAAAAGATATCCTCCAGCGCTTCAA ATACTATTTATCCAGAGGAGACGGACGCAACCGAGCACAGCTGGTAGCAGAATTAGACACGCTTTTGAAAGAAGCCCTCAG GTCGTATCCTGAGAAACATGGCCCAACAGATTAACACTTGTCACCTTGCCGCCAGTGAAAGCCTGATCTTAGCGCAACTCTTT TCAAAAATTGCAGGTCAACTGAGCCAGACCAATCCTGCTTCTGATTTGCTAGATGAAATTGAACGTTATCTGGAAGTCTTCCG GAACCGCAGTCTGCCCAAGACAAGAGAAGTTTGAAACCCGCGCCACCCTTCTTCAACTCCTACGTGAAGCCAAAACCTTC ATCCAAGTAAAAGTTGATTTTTACCAAAAATATAGACAGTAA

4158.6 (SEQ. ID. NO. 277)

ATGGAAATCATGTCGCTTGCGATTGCTGTTTTTGCCGTCATCATTGGTTTAGTCATTGGATATGTCAGCATCTCAGCTAAGATG AAATCATCTCAGGAAGCTGCAGAGTTGATGCTTTTAAATGCTGAACAAGAAGCAACTAATTTACGTGGACAAGCTGAGCGTG AAGCGGATTTACTTGTTAATGAAGCCAAACGTGAAAGCAAGTCTCTTAAAAAAGAAGCACTATTGGAGGCCAAAGAAGAAG AGAGAGCTACTAGCCTTGACCGTAAGGACGACAATTTGACGAGTAAAGAACAACACTTGAACAAAAAGAACAAAGTATTT CTGATAGAGCGAAAAACCTTGATGCGCGTGAAGAGCAATTAGAGGAAGTCGAAAGAACAAAAAGAAGCAGAACTAGAGCGT GCATTCGCGAAGCTGAGCAAGAGGTCAAGGAACGTTCTGACAAAATGGCCAAGGACATCTTGGTTCAAGCTATGCAACGTAT CGCTGGTGAATATGTAGCGGAGTCAACAAACTCAACAGTTCATCTGCCAGACGATACTATGAAGGGACGCATTATTGGTCGT GAAGGTCGTAACATTCGTACCTTTGAAAGTTTGACAGGGGTCGATGTGATTATCGACGATACACCAGAAGTGGTGACCTTGT CAGGATTTGATCCGATTCGTCGTGAGATTGCCCGTATGACTATGGAAAATGTTGCTCAAAGATGGTCGTATACATCCAGCTCGT ATCGAAGAGTTGGTTGAGAAAAACCGTCAAGAGATTGACAATAAGATTCGTGAATACGGTGAGGCTGCTGCCTATGAAATTG GTGCGCCAAACCTTCATCCAGACTTGATGAAGATTATGGGACGTTTGCAGTTCCGTACTTCATATGGACAAAATGTTTTGCGC CCTTCACGATATCGGGAAAGCCATTGACCATGAGGTTGAAGGTAGCCACGTTGAAATCGGTATGGAATTGGCCCGTAAGTAC AAGGAACCCCCAGTTGTGGTGAATACGATTGCTAGTCACCACGGAGATGTTGAAGCTGAGAGCGTGATAGCAGTTATCGTCG CTGCAGCAGATGCCTTGAGCGCAGCCCGTCCAGGTGCTCGTAGTGAGTCTCTTGAAAGCTACATCAAGCGTCTCCATGATTTG GAAGAAATTGCTAACGGCTTTGAAGGAGTGCAAACTAGCTTTGCCCTTCAAGCAGGACGTGAAATTCGTATCATGGTCAATC CAGGAAAAATCAAGGACGACAAAGTCACAATCTTGGCTCACAAAGTTCGTAAGAAAAATTGAAAACAATCTCGATTATCCAG GAAATATCAAGGTAACCGTGATTCGCGAGCTTCGTGCAGTAGATTATGCTAAATAA

4158.7 (SEQ. ID. NO. 278)
ATGATGTTAAAACCCTCTATTGATACCTTGCTCGACAAGGTTCCTTCAAAATATTCACTCGTAATCTTGGAAGCAAAACGTGC CCACGAATTGGAAGCAGGTGCCCCAGCAACTCAAGGTTTCAAGTCTGAAAAATCAACTCTTCGCGCTTTAGAAGAAATCGAA TCAGGAAACGTTACAATTCACCCAGATCCAGAAGGAAAACGTGAAGCAGTGCGTCGCCGTATCGAAGAAGAAAACGCCGC AAAGAAGAAGAAGAAAATCAAAGAGCAAATTGCTAAAGAAAAAGAAGATGGTGAAAAAATTTAA

4161.1 (SEQ. ID. NO. 279)

ATGTCAGCATATCAATTACCGACCGTATGGCAGGATGAAGCTAGTAATCAAGGAGCTTTTACGGGGCTAAACAGACCAACAG CAGGTGCCGTTTCGAACAAACTTGCCAAAAGGAGACAAGCTTTTCAGCTTTATTCACTGGGAACACCAAATGGTGTGAA GGTTACTATCTTATTGGAAGAATTACTAGAAGCTGGTTTTAAGGAAGCGGCTTACGACTTGTATAAGATTGCTATCATGGATG GGGATCAATTCGGATCAGACTTTGTGAAGCTCAATCCAAATTCCAAGATTCCAGCCTTATTGGACCAGTCAGGTACTGAAAA CGTAAGAGTCTTTGAGTCTGCTCATATTCTTCTTTACCTTGCTGAGAAATTTGGAGCCTTTTTACCAAGTAATCCTGTGGAAAA GGTAGAAGTTTTGAATTGGCTATTCTGGCAAGCAGGTGCAGCACCTTTTCTAGGTGGGGGATTTTGGACATTTCTTCAATTATG CTCCTGAAAAATTGGAATATCCTATTAACCGTTTTACGATGGAAGTGAAACGCCAGTTGGATTATTGGATAAGGAATTGGCT ATCTTTACCAAGGTTCTGCAAAATTCTTGGATGCCTCAAGTTATCAAAATCTAGTAAAATGGGCAGAAAAAATTGCCAATCG TCCAGCTGTTAAGCGTGGCTTGGAAGTAACTTATACAGAAATTAAATAG

4161.2 (SEO. ID. NO. 280)

TTGGCAAGCTTGATCACTTCTATCATCATGTTCTATGTCGGTTTCGATGTTCTAAGAGATACCATTCAAAAGATTCTCAGTCGG GAAGAAACGGTCATTGATCCTCTTGGTGCAACTCTAGGAATCATTTCTGCAGCGATTATGTTTGTGGTCTATCTCTACAATAC TCGCCTCAGTAAGAAATCCAACTCCAATGCGCTGAAGGCAGCTGCTAAGGACAATCTTTCTGACGCTGTTACCTCACTTGGA ACCGCCATTGCCATCCTAGCTAGTTTCAATTATCCGATTGTGGATAAACTGGTTGCTATCATCATCACTCTTTTATCTTG AAGACTGCCTATGATATCTTCATCGAGTCTTCCTTTAGTCTTTCAGATGGCTTTGACGACCGCCTGCTCGAGGACTACCAAAA GGCTATCATGGAAATTCCCAAAATCAGCAAGGTCAAATCGCAAAGAGGTCGCACCTACGGTAGCAACATCTACCTGGATATT ACACTAGAGATGAATCCTGACTTGTCTGTTTTTTGAAAGCCATGAAATCGCGGATCAGGTCGAGTCTATGCTGGAGGAGCGTT TTGGCGTCTTTGATACCGATGTCCATATCGAACCAGCACCTATCCCTGAGGATGAAATTTTTAGACAATGTCTATAAAAAATTG CTTATGCGTGAACAATTGATTGACCAAGGAAACCAACTAGAAGAACTCTTGACTGATGATTTTGTCTATATTCGCCAAGATG GAGAGCAGATGGATAAAGAGGCTTATAAGACCAAAAAAGAGTTAAATTCTGCTATCAAGGACATTCAAATTACTTCCATCAG ${\tt TCAAAAAACCAAACTCATCTGCTATGAGTTAGATGGTATCATCCATACCAGTATCTGGCGTCGCCACGAAACCTGGCAAAATATCTTTCATCAAGAAACCAAAAAAGAATAG}$

4162.1 (SEQ. ID. NO. 281)

4162.2 (SEQ. ID. NO. 282)

4164.2 (SEQ. ID. NO. 283)

ATGGAAAGTTTACTTATTCTATTATTAATTGCCAATCTAGCTGGTCTCTTTCTGATTTGGCAAAGGCAGGATAGGCAGGAGAA ACACTTAAGTAAGAGCTTGGAGGATCAGGCAGATCATTTGTCAGACCAGTTGGATTACCGCTTTGACCAAGCCAGACAAGCC AGCCAGTTAGACCAAAAAGATTTGGAAGTGGTTGTCAGCGACCGTTTGCAAGAAGTGCGGATTGAATTGCACCAAGGTCTGA AAATGAGCAACGTTTGGAACAATGCGCCAGACGGTCGAGGAAAAACTAGAAAAGACCTTGCAGACACGCTTACAGGCTTC CTTAACAAGGTTCTCTCTGGAACCAAGACGCGAGGGATTCTGGGGAGAATTGCAACTGGGGCAAATTATTGAAGACATCATGA CACCTGCCCAGTACGAACGAGAATACGCAACGGTTGAAAACTCTAGTGAACGAGTGGAGTATGCCATCAAGTTACCCGGAC AAGGCGACCAAGAATACGTCTATCTGCCAATTGACTCTAAGTTTCCACTGGCAGATTATTACCGCTTGGAAGAAGCCTATGA GACAGGTGACAAGGATGAGATTGAACGCTGTCGTAAGTCACTCCTAGCAAGCGTCAAGCGCTTTGCTAGGGATATTAGGAAC AAGTACATAGCACCACCTCGGACGACCAATTTTGGAGTTTTGTTTCTTCCGACAGAAGGTCTCTACTCAGAAATCGTCCGCAA TCCGGTCTTCTTTGATGATTTGAGACGGGAAGAACAGATTATTGTTGCAGGACCAAGTACCCTATCAGCCCTTCTTAACTCCC TATCAGTTGGTTTCAAGACCCTTAATATCCAAAAGAGTGCCGACCATATCAGCAAGACTCTTGCCAGTGTCAAGACCGAGTTT GGCAAGTTTGGTGGTATTCTGGTCAAGGCACAAAAACATCTCCAACATGCCTCTGGCAATATTGATGAATTATTAAACCGTC GTACCATAGCTATCGAGCGGACGCTCCGTCACATTGAGTTGTCAGAAGGTGAGCCTGCGCTTGATCTACTCCATTTTCAAGAA AATGAGGAAGATATGAAGATTAG

4164.3 (SEQ. ID. NO. 284)

4166.2 (SEQ. ID. NO. 285)

4166.3 (SEQ. ID. NO. 286)

ATGAAAATAGATATAACAAATCAAGTTAAAGATGAATTTCTTATATCATTAAAAACCTTGATTTCCTATCCTTCAGTACTCAA TGAAGGAGAAAATGGAACACCTTTTGGACAAGCAATCCAAGATGTCCTAGAAAAAACTTTAGAGATTTGTCGAGACATAGG TTTCACTACCTATCTTGACCCTAAAGGTTATTACGGATATGCAGAAATCGGTCAGGGAGCAGAGCTTCTGGCCATTCTCTGTC ATTTGGATGTTGTTCCATCAGGTGATGAAGCAGATTGGCAGACACCGCCATTTGAAGCAACTATCAAAGACGGCTGGGTATT CGGACGTGGTGTCCAAGATGATAAAGGCCCTTCGCTCGCAGCTCTCTATGCAGTAAAAAGCTTGCTGGACCAAGGTATTCAG TTCAAAAAGCGCGTACGCTTTATCTTTGGTACCGATGAGGAAACCCTCTGGCGCTGCATGGCACGCTACAATACCATCGAAG AACAGGCCAGTATGGGCTTTGCACCTGACTCATCTTTTCCTCTGACCTATGCTGAAAAAAGGGCTTCTACAGGTCAAACTTCAT GGCCCTGGATCGGATCAACTAGAGCTTGAAGTAGGAGGCGCCTTTAACGTTGTACCAGACAAGGCCAACTACCAAGGTCTCC TCTATGAACAGGTTTGTAACGGTCTCAAAGAAGCTGGTTATGATTACCAAACCACTGAACAAACCGTAACGGTTCTCGGAGT GCCAAAGCATGCTAAGGATGCTAGTCAAGGTATCAATGCTGTCATCCGACTAGCTACCATTCTTGCTCCTCCCAAGAACACC CTGCTCTCAGTTTTCTTGCAACACAAGCAGGTCAAGACGGCACAGGAAGACAAATCTTTGGTGATATAGCAGATGAACCTTC AGCTGACAAGGAAGAACTAGTAGAGTTGCTTACAAGATGTGCACAAAACTACCAACTCCGCTACGAAGAGTTTGACTATCTA GCGCCTCTATACGTCGCAGAAGACAGTAAACTCGTTAGCACACTGATGCAAATCTACCAAGAAAAGACTGGCGATAACAGTC CTGCTATTTCATCCGGTGGTGCCACTTTTGCTCGCACCATGCCAAATTGTGTAGCCTTCGGCGCCTTATTCCCAGGAGCGAAG CAGACAGAACATCAGGCAAATGAATGTGCCGTTCTAGAAGATTTGTACCGTGCTATGGATATTTATGCCGAAGCCGTCTATC GACTTGCAACTTAA

4169.1 (SEQ. ID. NO. 287)

ATGTCTAATTCATTTGTCAAGTTGTTAGTCTCCAATTATTTGCAAATTTAGCAGATATTTTCTTTAGAGTAACAATCATTGCT AACATATACATTATTTCAAAAATCAGTAATTGCCACATCACTAGTTCCTATCTTAATAGGAATATCCTCTTTTGTTGCGAGTCTT TTAGTTCCGTTGGTTACTAAAAGGTTAGCGCTAAATAGGGTTTTATCTTTATCTCAATTTGGAAAGACTATATTATTGGCGAT GTTTTGCAGCACCCGTTTCCTATGCTATTGTGCCACGCTATGCGACCGATTTGGGTAAGGCTAATTCAGCCTTATCAATGACT TTTAGTCTTGTATATCATTTCTAGCTTTCTGATGTTATTTCTTCCTAACGCTGAAGTGGAGGTGTTAGAGTCAGAAACTAATCT TGAAATTTTGCTCAAAGGTTGGAAGTTAGTTGCTAGAAATCCTAGATTAAGACTTTTTGTATCAGCAAATTTATTGGAAATTT CTAATACAGCATACTCTATTGGTATTATAATTAGTGGCTTAATTGCTTTTTAGGCTATCTGAAAAGTTCCTTGCTGCTAAATGGG AACCCCAATTATTCACCCCAAATCTAAAAACCATCCAGAATCCTTGCCTTAGCTTAGATCCTGGATGGTTTCTTTTTTCACCCA CATGAACAGTTTACCAAATCATCACTTCCAAAACAAGTCTTTTTACCAACTATCTTTCGATGGAGGTCATTTAACCCAGTATG GTGGTCTTATCTTTTTTCAGGAACTTTTTTCCCAGTTGAAACTAAAAGAGCGGATTTCTAAGTATTTAGTAACGAATGACCAA CGCCGCTACTGTCGTTATTCGGATTCAGATATCCTTGTCCAGTTCCTCTTTCAACTGTTAACAGGTTATGGAACGGACTATGCT TGTAAAGAATTGTCAGCTGATGCCTACTT

4169.3 (SEQ. ID. NO. 288)

4169.4 (SEQ. ID. NO. 289)

ATGCTTAAACGCTTATGGATGATCTTCGGACCGGTCTTGATCGCTGGTTTGTTGGTTTTTTCTGCTCATTTTCTTTTATCCTACTG AGATGCATCATAATCTAGGAGCTGAAAAGCGTTCAGCAGTGGCTACTACTATCGATAGTTTTAAGGAGCGAAGTCAAAAAGT CAGAGCACTATCTGATCCAAATGTGCGTTTTGTTCCCTTCTTTGGCTCTAGTGAATGGCTTCGTTTTTGACGGTGCTCATCCTGC GGTATTAGCTGAGAAATACAATCGTTCCTACCGTCCTTATCTTTTAGGACAGGGGGGAGCTGCATCGCTTAACCAATATTTTG GAATGCAACAGATGTTACCACAGCTGGAGAATAAACAAGTTGTGTATGTTATCTCACCTCAGTGGTTCAGTAAAAAATGGCTA TGATCCAGCAGCCTTCCAGCAGTATTTTAATGGAGACCAGTTGACTAGTTTTCTGAAACATCAATCTGGGGATCAGGCTAGTC AATATGCAGCGACTCGCTTACTGCAACAGTTCCCAAACGTAGCTATGAAGGACCTGGTTCAGAAGTTGGCAAGTAAAGAAGA ATTGTCGACAGCAGACAATGAATGAATTGAATTATTGGCTCGTTTTAATGAACGCCAAGCTTCCTTTTTTGGTCAGTTTTCGG TTAGAGGCTATGTTAACTACGATAAGCATGTAGCTAAGTATTTAAAAATCTTGCCAGACCAGTTTTCTTATCAGGCAATAGAA GATGTTGTCAAAGCAGATGCTGAAAAAAAATACTTCCAATAATGAGATGGGAATGGAAAATTATTTCTATAATGAGCAGATCA AGAAGGATTTGAAGAAATTAAAGGATTCTCAGAAAAGCTTTACCTATCTCAAGTCGCCAGAGTATAATGACTTGCAGTTGGT TTTAACACAGTTTTCTAAATCTAAGGTAAACCCGATTTTTATCATTCCACCTGTTAATAAAAAATGGATGAACTATGCTGGTC TACGAGAGGATATGTACCAACAAACGGTGCAGAAGATTCGCTACCAGTTAGAAAGTCAAGGTTTTACCAATATAGCAGATTT TTGATCCTTTCCTATCCAATCCCACACCAGCTCCGACTTACCATCTGAATGAGCGCTTTTTCAGCAAAGATTGGGCGACTTAT GATGGAGATGTCAAAGAATTTCAATAG

4169.6 (SEQ. ID. NO. 290)

4170.3 (SEQ. ID. NO. 291)

4170.4 (SEQ. ID. NO. 292)

ATGACTAATTTAATTGCAACTTTTCAGGATCGTTTTAGTGATTGGTTGACAGCTCTATCTCAACATTTGCAGTTGTCGCTTTTG ACCTTGTTACTAGCTATTTTGCTTGCGATTCCCTTGGCTGTTTTTCTTCGCTATCATGAGAAGCTGGCCGACTGGGTCTTGCAG ATTGCAGGTATTTTCCAGACCATCCCGTCTCTGGCCTTGTTGGGGCTCTTTATCCCTTTGATGGGAATTGGGACCTTGCCGGCT TTGACAGCTCTAGTGATTTATGCGATTTTCCCTATTTTGCAAAATACTATCACTGGGCTGAAGGGAATTGATCCGAACCTGCA AGAGGCTGGGATTGCCTTTGGGATGACCAGATGGGAACGTCTCAAGAAATTTGAAATTCCACTCGCCATGCCTGTTATCATG TCTGGGATTCGGACGGCAGCTGTTTTGATTATCGGTACGGCAACCTTGGCGGGCCTTGATTGGTGCAGGGGGACTAGGTTCCTT TATTCTTTTGGGAATTGACCGTAATAATGCCAGTTTGATTTTGATTGGGGCACTTTCTTCTGCAGTGCTAGCCATTGCCTTTAA CTTCCTACTAAAAGTGATGGAAAAAGCAAAATTACGGACGATTTTCTCAGGTTTTGCCTTGGTGGCTTTATTACTGGGTCTGT CTTATAGTCCAGCTCTTTTGGTTCAAAAAGAGAAGGAAAACTTGGTTATTGCTGGGAAAATAGGTCCAGAACCAGAAATTTT GGCCAATATGTATAAGTTGCTGATTGAAGAAAATACCAGCATGACTGCGACTGTTAAACCGAATTTTGGGAAGACAAGCTTC CTTTATGAAGCTCTGAAAAAAGGCGATATTGACATCTATCCTGAATTTACTGGTACGGTGACTGAAAGTTTGCTTCAACCATC ACCCAAGGTGAGTCATGAACCAGAACAGGTTTATCAGGTGGCGCGTGATGGCATTGCTAAGCAGGATCATCTAGCCTATCTC AAACCCATGTCTTATCAAAACACCTATGCTGTAGCTGTTCCGAAAAAGATTGCTCAAGAATATGGCTTGAAGACCATTTCAG ACTTGAAAAAAGTGGAAGGCAGTTGAAGGCAGGTTTTACACTCGAGTTTAACGACCGTGAAGATGGAAATAAGGGCTTGC ATCACGGATGCCTATTCGACTGATGCGGAATTGGAGCGTTATGATTTACAGGTCTTGGAAGATGACAAGCAACTCTTCCCAC CTTATCAAGGGGCTCCACTCATGAAAGAAGCTCTTCTCAAGAAACACCCAGAGTTGGAAAGAGTTCTTAATACATTGGCTGG TAAGATTACAGAAAGCCAGATGAGCCAGCTCAACTACCAAGTCGGTGTTGAAGGCAAGTCAGCAAAGCAAGTAGCCAAGGA GTTTCTCCAAGAACAAGGTTTGTTGAAGAAATGA

4170.5 (SEQ. ID. NO. 293)

4170.6 (SEQ. ID. NO. 294)

ATGAGAATATGAAGGCAAAATATGCTGTTTGGGTGGCTTTTTTCTTAAATTTGACTTATGCCATTGTTGAGTTTATTGCAGG
TGGAGTATTTGGTTCTAGCGCTGTTCTTGCTGACTCTGCATGACTTGGGAGATGCGATTGCAATTGGAATATCAGCTTTTCT
AGAAACAATCTCCAATCGTGAAGAAGACAATCAGTACACCTTGGGCTATAAGCGGTTTAGCCTGCTAGGAGCCTTGGTAACA
GCTGTGATTCTCGTAACGGGCTCTGTTCTAGTCATTTTGGAAAATGTCACGAAGATTTTGCATCCGCAACCAGTCAATGATGA
GGGGATTCTCTGGTTAGGAATTATTGCGATTACTATCAATCTGTTAGCGAGTCTGGTGGTTAGGGAAAGGAAAAGAAT
GAGTCTATTCTGAGTCTGCATTTTCTGGAAGATACCTTAGCGGTGGTTATCCTGATGGCGATTGTTCTTCGATTTAC

4170.8 (SEO. ID. NO. 295)

ATGATTGAATACAAAAATGTAGCACTGCGCTACACAGAAAAGGATGTCTTGAGAGATGTCAACTTACAGATTGAGGATGGG
GAATTTATGGTTTTAGTAGGGCCTTCTGGGTCAGGTAAGACGACCATGCTCAAGATGATTAACCGTCTTTTGGAACCAACTGA
TGGAAATATTTATATGGATGGGAAGCGCATCAAAGACTATGATGAGCGTGAACTTCGTCTTTTCTACTGGTTATGTTTTACAGG
CTATTGCTCTTTTTCCAAATCTAACAGTTGCGGAAAATATTGCTCTCATTCCTGAAATGAAGGGGTGGAGCAAGGAAGAATT
ACGAAGAAAACAGAAGAGCTTTTGGCTAAGGTTGGTTTACCAGTAGCCGAGTATGGCATCGCTTACCTAGTGAATTATCTG
GTGGAGACAGCAACGGGTCGGTATTGTCCGAGCTATGATTGGTCAGCCCAAGATTTTCCTCATGGATGAACCCTTTTCGGCC
TTGGATGCTATTTCGAGAAAACAGTTGCAGGTTCTGACAAAAGAATTGCATAAAGAGTTTTGGAACAACGATTTTTGTAA
CCCATGATACCGATGAAGCCTTGAAGTTGGCGGACCGTATTGCTCTTTGCAGGATGAGAAATTCGCCAGGTAGCGAATCC
CGAGACAATTTTAAAAGCGCCTGCAACAGACTTTTGTAGCAGACTTTTTGGAGGTAGTTCATGACTAA

4171.1 (SEQ. ID. NO. 296)

4172.1 (SEQ. ID. NO. 297)

TTGGTCATTTTACTAGGCTCTCTTTTTGAGCTTGCCCTTTGTCCAAGTTGAAAGCTCACGAGCGACTTATTTTGATCATCTTT TCACTGCTGTCTCCAGTCTGTGACGGGTCTCTCAACCCTTCCAGTAGCTCACACCTATAATATCTGGGGTCAAATAATC TCTTCGTAGCCGTGCAACTATTCAGGATAGTTTTAGTTATGGAGAAACTCGATCTTTGAGAAAGTTTGTCTATTCTATTTTTCT CACGACCTTTTTGGTTGAGAGCTTGGGAGCTATTTTGCTTAGTTTTCCCCTTATTCCTCAACTTGGCTGGGGACGTGGTCTTTT TAGTTCCATTTTTCTAGCGATCTCAGCCTTCTGTAATGCCGGTTTTGATAATTTAGGGAGCACCAGTTTATTTGCTTTTCAGAC CGATTTACTGGTCAATCTGGTGATTGCAGGCTTGATTATTACAGGCGGCCTTGGTTTTATGGTCTGGTTTGATTTGGCTGGTCA TGTAGGAAGAAAAAAGGACGTCTGCACTTTCATACGAAGCTTGTACTATTATTGACTATAGGTTTGTTGTTATTTGGAA TTTCAAACAGTGACGATGCGAACAGCTGGCTTTTCTACGATAGATTATACTCAGGCTCATCCTGTGACTCTTTTGATTTATATC TTACAGATGTTTCTAGGTGGGGCACCTGGAGGAACAGCTGGGGGACTCAAGATTACGACATTTTTTTGTCCTCTTTGGTCTTTGC CTCGTATTTGAAACCATTTCAGCTCTTAGTACAGTTGGTGTAACGGCAAATCTGACTCCTGACCTTGGGAAATTGGCTCTCAG **TGTTATCATGCCACTTATGTTTATGGG**

4172.2 (SEQ. ID. NO. 298)

4172.3 (SEO, ID, NO, 299)

4172.4 (SEQ. ID. NO. 300)

ATGAAATTCAATCCAAATCAAAGATATACTCGTTGGTCTATTCGCCGTCTCAGTGTCGGTGTTGCCTCAGTTGTTGTGGCTAG TGGCTTCTTTGTCCTAGTTGGTCAGCCAAGTTCTGTACGTGCCGATGGGCTCAATCCAACCCCAGGTCAAGTCTTACCTGAAG CTGGAAATACGAATTCACTTCCGACACCTACAGAAAGAACTGAAGTGAGCGAGGAAACAAGCCCTTCTAGTCTGGATACACT TTTTGAAAAAGATGAAGAAGCTCAAAAAAATCCAGAGCTAACAGATGTCTTAAAAGAAACTGTAGATACAGCTGATGTGGA ATGTTCCTGCTGCTTATCTTGAAAAAGCTGAAGGGAAAGGTCCTTTCACTGCCGGTGTAAACCAAGTAATTCCTTATGAACTA TTCGCTGGTGATGGTATGTTAACTCGTCTATTACTAAAAGCTTCGGATAATGCTCCTTGGTCTGACAATGGTACTGCTAAAAA TCCTGCTTTACCTCCTCTTGAAGGATTAACAAAAGGGAAATACTTCTATGAAGTAGACTTAAATGGCAATACTGTTGGTAAAC AAGGTCAAGCTTTAATTGATCAACTTCGCGCTAATGGTACTCAAACTTATAAAGCTACTGTTAAAGTTTACGGAAATAAAGA CGGTAAAGCTGACTTGACTAATCTAGTTGCTACTAAAAATGTAGACATCAACATCAATGGATTAGTTGCTAAAGAAACAGTT TTCACAGCAGGTGTCAACCATGTGATTCCATACGAACTCTTCGCAGGTGATGGCATGTTGACTCGTCTCTTGCTCAAGGCATC TGACAAGGCACCATGGTCAGATAACGGCGACGCTAAAAACCCAGCCCTATCTCCACTAGGCGAAAACGTGAAGACCAAAGG TCAATACTTCTATCAAGTAGCCTTGGACGGAAATGTAGCTGGCAAAGAAAAACAAGCGCTCATTGACCAGTTCCGAGCAAAT GGTACTCAAACTTACAGCGCTACAGTCAATGTCTATGGTAACAAAGACGGTAAACCAGACTTGGACAACATCGTAGCAACTA AAAAAGTCACTATTAACATAAACGGTTTAATTTCTAAAGAAACAGTTCAAAAAAGCCGTTGCAGACAACGTTAAAGACAGTAT CGATGTTCCAGCAGCCTACCTAGAAAAAGCCAAGGGTGAAGGTCCATTCACAGCAGGTGTCAACCATGTGATTCCATACGAA CTCTTCGCAGGTGATGGTATGTTGACTCGTCTCTTGCTCAAGGCATCTGACAAGGCACCATGGTCAGATAACGGTGACGCTAA AAACCCAGCCCTATCTCCACTAGGTGAAAACGTGAAGACCAAAGGTCAATACTTCTATCAATTAGCCTTGGACGGAAATGTA GCTGGCAAAGAAAAACAAGCGCTCATTGACCAGTTCCGAGCAAACGGTACTCAAACTTACAGCGCTACAGTCAATGTCTATG GTAACAAGACGGTAAACCAGACTTGGACAACATCGTAGCAACTAAAAAAAGTCACTATTAACATAAACGGTTTAATTTCTAA

4172.5 (SEQ. ID. NO. 301)

4172.6 (SEQ. ID. NO. 302)

4174.1 (SEQ. ID. NO. 303)

4175.2 (SEQ. ID. NO. 304)

4175.3 (SEQ. ID. NO. 305)

GGAAGGTCGTGTTGTCCTTTCTGGTGGTCCAGAATTGGCTGCGCGTTTTGGAACGTGAAGGATACGCAAAATTAGCTGAAGAA CTTGGCTACGACTACAAGGAAGAATTGTAA

4174.4 (SEQ. ID. NO. 306)

ATGCCCTACAAAAGACAAAGGAGTTTTTCAATGGCACTTTCTAAACTAGATAGCCTTTATATTGGCAGTGGTAGCAGACCATT CGAAAAATCCACATCACCAAGGGAAGTTAGAAGATGCTGAGCAAATCAGTCTCAACAATCCGACTTGTGGGGATGTCATCA ACCTCTCTGTCAAGTTTGATGCAGAGGACCGTTTTGGAAGATATTGCTTTTCTAAATTCAGGATGCACGATTTCAACTGCTTCT GCTAGTATGATGACAGATGCCGTTTTAGGAAAAACCAAACAAGAAATTTTAGAACTGGCGACTATTTTTTTCTGAAATGGTTC AAGGGCAAAAAAGATGAGGTCAAGACCAACTTGGAGACGCGGCATTCTTGTCAGGTGTTGCCAAATTCCCTCAAAGAATCAAGTGTGCAACCCTAGCTTGGAATGCCCTTAAGAAAACAATTGAAAATCAAGAAAAACAGTAA

4175.5 (SEQ. ID. NO. 307)

ATGAAAATTCAAGACCTATTGAGAAAAGATGTCATGTTGCTAGATTTGCAGGCAACTGAAAAAACAGCTGTCATCGACGAGA TGATTAAAAATTTGACAGACCACGGTTATGTAACAGATTTTGAAACATTTAAAGAAGGAATTTTGGCGCGTGAAGCTTTGAC TTCTACTGGTTTGGGTGATGGAATCGCAATGCCTCACAGCAAAAACGCTGCTGTCAAAGAAGCGACAGTTCTATTTGCTAAG TCAAATAAGGGTGTTGACTACGAGAGCTTGGATGGACAAGCAACTGACCTCTTCTTCATGATTGCAGCTCCAGAAGGTGCCA ATGATACTCACTTGGCAGCCTTGGCAGAATTGTCTCAATACTTGATGAAAGACGGTTTTGCAGACAAACTTCGTCAAGCAAC ATCTGCAGACCAAGTTATCGAACTTTTTGACCAAGCTTCAGAAAAAACTGAGGAACTTGTTCAAGCACCTGCTAATGACTCT GGTGACTTTATCGTAGCTGTTACAGCTTGTACAACAGGTATTGCCCACACTTACATGGCCCAAGAAGCCCTTCAAAAAGTAG TAAGGCTAAAGCTATTATCATTGCAGCAGACAAGGCCGTTGAAATGGATCGATTTGATGGAAAACCATTGATCAATCGTCCA GTTGCTGACGGTATCCGTAAGACAGAAGAGCTAATTAACTTGGCTCTTTCAGGAGATACTGAAGTCTACCGTGCCGCTAATG GTGCCAAAGCTGCAACAGCCTCTAACGAAAAACAAAGCCTTGGTGGTGCCTTGTACAAACACTTGATGAGTGGTGTATCTCA CCTTGGCAATCTTGGTTCTTACCATGAGTTAGCTTCTATGTTCATGAAAATTGGTGGAGCTGCCTTTGGTTTGATGCTTCCAGT CTTTGCGGGTTATGTTGCCTACTCTATTGCTGAAAAACCGGGTTTGGTAGCAGGTTTCGTGGCTGGTGCTATTGCCAAAGAAG GTTTTGCCTTTGGTAAAATTCCTTATGCCGCAGGTGGTGAAGCAACTTCAACTCTTGCAGGTGTCTCATCTGGTTTCCTAGGTG CCCTTGTTGGTGGATTTATCGCAGGTGCCTTGGTTCTTGCCATCAAGAAATACGTTAAAGTTCCTCGTTCACTCGAAGGTGCT AAATCAATCCTTCTATTGCCACTTCTTGGAACAATCTTGACAGGATTTGTTATGCTAGCTGTGAATATCCCAATGGCTGCAAT CAACACTGCTATGAATGACTTCCTAGGCGGTCTTGGAGGAGGTTCAGCTGTCCTTCGTAGTCGTCCTTGGTGGAATGATGG CTGTTGACATGGGTGGACCAGTTAATAAAGCAGCTTATGTCTTTTGGTACAGGTACGCTTGCAGCAACTGTTTCTTCAGGTGGT TCTGTAGCCATGGCAGCAGTTATGGCTGGAGGAATGGTGCCACCACTTGCAATCTTTGTCGCAACTCTTCTTTTCAAAGATAA ATTTACTAAGGAAGAACGTAACTCTGGTTTGACAAACATCATCATGGGCTTGTCATTTATCACTGAGGGAGCGATTCCATTTG GTGCCGCTGACCCAGCTCGTGCGATTCCAAGCTTCATCCTTGGTTCAGCAGTAGCAGGTGGACTCGTTGGTCTTACTGGTATC AAACTCATGGCGCCACACGGAGGAATCTTCGTTATCGCCCTTACTTCAAATGCTCCTTTACCTCGTTTCTGTCTTGGTAGGA GCAATCGTAAGTGGTGTGGTTTATGGTTACCTACGCAAACCACAAGCATAA

4175.6 (SEQ. ID. NO. 308)

GTTGATTTCGTTAGGAATTGCGATTTTATTGATTTTCGCAGCCTTCAAATTAGGGGCTGCAGGTATAACCCTTTATAATTTAAT GGAAGGACTCTTATCTGGCTTTTTCACCATATTTGCTGGCTTACTCTTGATTTTTGAGGCCTACTTGGTTTGGAAATATGGTTT GGACAAGTCCGTTCTAAAAGGGACCATGGCTCAGGTTGTGACAGATCTGACTGGTTTTCGAACGACTAGCTTTGCTGGAGGG TAGTGGGTTCTCCTAGTCAGCCCTTGGTCTGTTTACGATATTGCTGAATTTTTCAGTAGAGGCTTTGCCAAATGGTGGGAA GGGCACGAGCGTCGAAAAGAGGAACGCTTTGTCAAACAAGAAGAAAAAGCTCGCCAAAAGGCTGAGAAAGAGGCTAGATT AGAACAAGAAGAGACTGAAAAAGCCTTACTCGATTTGCCTCCTGTTGATATGGAAACGGGTGAAATTCTGACAGAGGAAGC GAACAGGAAGATGACTCAGATGACGAAGATGTTCAGGTCGATTTTTCAGCCAAAGAAGCCCTTGAATACAAACTTCCAAGCT TACAACTCTTTGCACCAGATAAACCAAAAGATCAGTCTAAAGAGAAGAAAATTGTCAGAGAAAATATCAAAAATCTTAGAAG CAACCTTTGCTAGCTTTGGTATTAAGGTAACAGTTGAACGGGCCGAAATTGGGCCATCAGTGACCAAGTATGAAGTCAAGCC GGCTGTTGGTGTAAGGGTCAACCGCATTTCCAATCTATCAGATGACCTCGCTCTAGCCTTGGCTGCCAAAGATGTCCGGATTG AAGCACCAATCCCTGGGAAATCCCTAATCGGAATTGAAGTGCCCAACTCCGATATTGCCACTGTATCTTTCCGAGAACTATG GGAACAATCGCAAACGAAAGCAGAAAATTTCTTGGAAATTCCTTTAGGGAAGGCTGTTAATGGAACCGCAAGAGCTTTTGAC CTTTCTAAAATGCCCCACTTGCTAGTTGCAGGTTCAACGGGTTCAGGGAAGTCAGTAGCAGTTAACGGCATTATTGCTAGCAT TCTCATGAAGGCGAGACCAGATCAAGTTAAATTTATGATGGTCGATCCCAAGATGGTTGAGTTATCTGTTTACAATGATATTC CCCACCTCTTGATTCCAGTCGTGACCAATCCACGCAAAGCCAGCAAGGCTCTGCAAAAGGTTGTGGATGAAATGGAAAACCG TTATGAACTCTTTGCCAAGGTGGGAGTTCGGAATATTGCAGGTTTTAATGCCAAGGTAGAAGAGTTCAATTCCCAGTCTGAGT ACAAGCAAATTCCGCTACCATTCATTGTCGTGATTGTGGATGAGTTGGCTGACCTCATGATGGTGGCCAGCAAGGAAGTGGA AGATGCTATCATCCGTCTTGGGCAGAAGGCGCGTGCTGCAGGTATCCACATGATTCTTGCAACTCAGCGTCCATCTGTTGATG TCATCTCTGGTTTGATTAAGGCCAATGTTCCATCTCGTGTAGCATTTGCGGTTTCATCAGGAACAGACTCCCGTACGATTTTGG ATGAAAATGGAGCAGAAAAACTTCTTGGTCGAGGAGACATGCTCTTTAAACCGATTGATGAAAATCATCCAGTTCGTCTCCA AGGCTCCTTTATCTCGGATGACGATGTTGAGCGCATTGTGAACTTCATCAAGACTCAGGCAGATGCAGACTACGATGAGAGT TTTGATCCAGGTGAGGTTTCTGAAAATGAAGGAGAATTTTCGGATGGAGATGCTGGTGGTGATCCGCTTTTTGAAGAAGCTA AGTCTTTGGTTATCGAAACACAGAAAGCCAGTGCGTCTATGATTCAGCGTCGTTTATCAGTTGGATTTAACCGTGCGACCCGT CTCATGGAAGAACTGGAGATAGCAGGTGTCATCGGTCCAGCTGAAGGTACCAAACCTCGAAAAGTGTTACAACAATAA

4176.1 (SEQ. ID. NO. 309)

4178.2 (SEO, ID, NO, 310)

ATGAATAAATCAGAACACCGCCACCAACTTATACGCGCTCTTATCACAAAAAACAAGATTCATACACAGGCTGAGTTGCAAG
CCCTTCTTGCTGAGAACGACATTCAAGTAACCCAGGCAACCCTCTCACGCGACATCAAAAATATGAACCTATCAAAAGTCCG
CGAAGAAGATAGCGCTTATTATGTTCTTAACAATGGTTCCATCTCAAAATGGGAAAAACGTCTCGAACTCTACATGGAAGAC
GCCCTTGTCTGGGATGCGCCCAGTTCAACACCAAGTCCTACTAAAAACCCTTCCTGGACTGGCTCAATCCTTTTGGTTCTATCATT
GATACTTTGAGCTTCCCTGACGCTATCGCTACCCTTTGTGGTAATGATGTCTTTTTCATCTGTGAAGATGCAGATACTGCT
CAAAAGTGCTTTGAAGAACTGAAAAAATTCGCCCCCACCATTTTTCTTTTGAAGAATAA

4179.1 (SEQ. ID. NO. 311)

ATGAAAAGTATAAAATTAAATGCTCTATCTTACATGGGAATTCGTGTCTTGAATATTATTTTTCCCATCCTAACTGGAACCTA TGTCGCGCGTGTCTTGGACCGAACTGACTATGGTTACTTCAACTCAGTCGACACTATTTTGTCATTTTTCTTGCCCTTTTGCAAC TTATGGTGTCTATAACTACGGTTTAAGGGCTATCAGTAATGTCAAGGATAACAAAAAAAGATCTTAACAGAACCTTTTCTAGTC TTTTTTATTTGTGCATCGCTTGTACGATTTTGACCACTGCTGTCTATATCCTAGCCTATCCTCTTCTTTACTGATAATCCAAT CGTCAAAAAGGTCTACCTTGTTATGGGGATTCAACTCATTGCCCAGATTTTTTCAATCGAATGGGTCAATGAAGCTCTGGAAA ATTACAGTTTTCTCTTTTTACAAAACTGCCTTCATCCGTATCCTGATGCTGGTCTCTATTTTCTTATTTGTTAAAAAATGAACACG ATATTGTTGTCTATACACTTGTGATGAGTTTATCGACGCTGATTAACTACCTGATTATTTTTTGGATTAAAAAGAGACATCA AACTTGTTAAAATTCACCTAAGTGATTTTAAACCACTCTTTCTCCCTCTGACAGCCATGTTAGTCTTTTGCCAATGCCAATATGC AGAAGCCTATGTTTCTCTGGTTAATAGAGGTAGTCGAATCTTTAACTTCTTTATCATTCCACTGAGTTTTGGACTCATGGTTTT TTATCCTGGCCTTAGATACCATTCTTGGTTCCCAAATTCTCTTTACCAATGGCTATGAAAAACGTATCACAGTCTATACAGTCT TGCTATCAGAGACTTCTCTACTTGTTTTCTATATCATTTTCATCCATAGAAAACAACTCATCCACTTGGGACATATCTTTAGCT TAATTTGCCATTTTTGATTA

ATACTGGTTTGATTGTCTTGCTATCAGCTATCTCTTATATTAGTCTACTTGTCTTCACAAAAGATAGCATTTTCTATGAATTTTT AAACCATGTCCTAGCCTTAAAAAAATAAATTTAAAAAATCATAG

4179.2 (SEQ. ID. NO. 312)

4179.3 (SEQ. ID. NO. 313)

GCCAAATGGTGTTGGGAAATCAACTCTGTTAGCCAAGTTACAGAGACTTCTGAATGATAAAAGAGAGATTTCACTTGGTTTT
ATGCCACAAGATTACCACAAAAAACTGCAATTGGATTTATCCCCAATAGCCTATCTCAGTAAAACTGGGGAAAAAGAGGAA
CTACAGAAAATCCAATCTCACCTAGCTAGTCTCAATTTCAGTTATCCAGAAATGCAGCATCAAATTCGCTCCTTATCTGGCGG
ACAACAGGGAAAACTCCTGCTTTTGGATTTAGTCCTGCGCAAACCAAACTTTCTCCTGCTGGATGAACCCACACAGAAACTTTT
CTCCCACTTCTCAACCCCAAATCAGAAAACTCTTTGCTACCTATCCAGGCGGTCTCATCACTGTTTCGCATGACGTCGTTTTC
TAAAAGAAGTCTGCTCGATCATCTATCGCATGACAGAACACGGTTTTGAAAGCTAGTTAATTTAGAAGATTTTATAA

4179.4 (SEQ. ID. NO. 314)

4179.6 (SEQ. ID. NO. 315)



ATGAGTATTAAACTAATTGCCGTTGATATCGACGGAACCCTTGTCAACAGCCAAAAGGAAATCACTCCTGAAGTTTTTTCTGC
CATCCAAGATGCCAAAGAAGCTGGTGTCAAAGTCGTGATTGCAACTGGCCGCCCTATCGCAGGCGTTGCCAAACTTCTAGAC
GACTTGCAGTTGAGAGACGAGGGGGACTATGTGGTAACCTTCAACGGTGCCCTTGTCCAAGAAACTGCTACAGGACATGAGA
TTATCAGCGAATCCTTGACTTATGAGGATTATCTAGATATGGAATTCCTCAGTCGCAAGCTCGGTGTCCACATGCCATT
ACCAAGGACGGTATCTATACTGCAAATCGCAATATCGGAAAATACACTGTACACGAATCAACCCTCGTCAGCATGCCTATCT
TCTACCGTACCCCTGAAGAAATGCTTGACAAAAATTGTTAAATGTTATCGATGAACCAGAAATTCTCGATGCTGC
GATTGAAAAAATTCCAGCAGAATTTTACGAGCGCTACTCCATCAACAAATCTGCTCCTTTCTACCTCGAACTCCTTAAAAAGA
ATGTAGACAAGGGTTCAGCCATTACTCACTTGGCTGAAAAACTCGGATTGACCAAAGATGAAACCATGGCAATCGGTGATGA
AGAAAATGACCGTGCCATGCTGGAAGTCGTTGGAAACCCCGTTGTCATGGAAAAATCGAAAATCAAAAAAATCGC
CAAATACATCACCAAAACAAATGACGAATCCGGCGTTGCCCATCCCATCCGAACATGGGTACTGTAA

4179.7 (SEQ. ID. NO. 316)

4179.8 (SEQ. ID. NO. 317)

ATGTTGTTTGATCAAATTGCAAGCAATAAACGAAAAACCTGGATTTTGTTGCTGGTATTTTTCCTACTCTTAGCTCTTGTTGGT
TATGCGGTTGGTTATCTCTTTATAAGATCTGGACTTGGTGGTTTTGGTTATTGCACTGATTATCGGCTTTATCTACGCTTTGTCT
ATGATTTTTCAATCGACAGAGATTGTCATGTCCATGAATGGAGCGCGTGAGGTGGATGAGCAAACGGCACCAGACCTCTACC
ATGTAGTGGAAGATATGGCTCTGGTCGCTCAGATTCCTATGCCCCGTGTTTTCATCATTGATGATCCAGCCTTAAATGCCTTTG
CGACAGGTTCTAATCCTCAAAATGCGGCTGTTGCTGCGACTTCAGGTCTACTAGCTATCATGAATCGTGAAGAACCTAGAAGC
TGTTATGGGACATGAAGTCATATTCGTAATTATGATATCCGTATTTCGACTATTGCAGTTGCCCTTGCTAGTGCTATCAC
CATGCTTTCTAGTATGGCAGGTCGTATGATGTGGTGGGGTGGAGCAGGTCGCAGACGAAGTGATGACCGAGATGAAAT
GGTCTTGAAATCATTATGCTAGTGGTTTCCCTACTAGCTATTGTACTGGCACCTCTCCTGCAACCTTTGGTTCAGCTCGCTATT
TCTCGTCAGAGGGAATTTCTGGCAGATGCATCTAGTGTCGAGCTGACTCGCAATCCCCAGGGAATGATTAATGCCTAGATA
AGTTGGACAATAGCAAACCTATGAGTCGCCACCTCGATGATCCCAGAGAATGATCCAAAAAAGGTGG
GGGGTTCCAAAAACCTCTTTTATACCCACCCACCTATCTCAGAACGATTGAACGTTTAAAACAGATGTAA

4179.9 (SEQ. ID. NO. 318)

4179.12 (SEO. ID. NO. 319)

4181.1 (SEQ. ID. NO. 320)

ATGAACGTCAATTAGCCTTGGTCGTCTTTAGTGGTGGTCAAGATTCAACAACCTGCCTTTTCTGGGTCATGCAACACTATGA
AACAGTCGAAGCTGTCACCTTTGCCTACGGCCAACGTCATCACCTCGAAATTCAAATTACTAGAGAAATCGCTAAGGAACAG
GGCATTCGTCACCCATATCCTCGATATGTCTCTGCTGGGACAAATCACTGCTCAGCCAGACTTTGCGACGAGATTCATATTTCCTA
CATTCCTGACAAGCTCTGTGGGGAAAATCCCTCAAACTATTATCTATTTAGCTACCGGAAACCACGGAGATTCCACGCAAA
ACTGTATCAACACCACCGGGAAAGCTTTGGTCAACTTGCTAGACCCTCGCTATTTAGAAGTCTGGGGAAAAATTCACTCCGCG
CGGTGGCATTTCAATCGACCCCTACTACAACTACGGTAAGCAAGGAACTAAGTATGAGGGCTTGGCAGAACAACGCCTCTTC
CAACACGACCTTTATCCAGAGAAAATTGACAACCGCTAA

4181.2 (SEQ. ID. NO. 321)

ATGACCGAAACGGTAGAAGATAAAGTAAGTCATTCAATTACTGGGCTTGATATCCTCAAGGGGATAGTTGCTGCGGGAGCTG
TCATAAGTGGAACCGTTGCAACTCAAACGAAGGTATTTACAAATGAGTCAGCAGTACTTGAAAAAAACTGTAGAGAAAACCG
ATGCTTTGGCAACAAATGATACAGTAGTTCTAGGTACGATATCTACAAGTAATTCAGCGAGTTCAACTAGTTTGTCAGCTTCA
GAGTCGGCAAGTACATCTGCATCTGAGTCAGCCTCAACCAGCGCTTCGACCTCAGCAAGTACAAGTGCATCAGAATCAGCAA
GTACATCGGCTTCGACAAGTATTTCTGCATCATCTACTGTGGTAGGTTCACAAAACAGCTGCCGCTACAGAAGCAACTGCTAA
GAAGGTCGAAGAAGAACCAGCTAGTGATTATGTAAGCATCAGTTACAAATGTCAATCTCAATCTTATGCTAAG
CGACGCAAGCGTTCAGTGGATTCCATCGAGCAATTGCTGCTTTATAAAAAATGCTGCTGTTTTTTCTGGCAATACGATTGT
AAATGGCGCCCCTGCAATTAATGCAAGTCTAAAACATTGCTAAAAGTGAGACAAAAAGTTTAACAGGTGAAGGTGTAGGTTCG
GTATATCGTGTTCCAATTTACTATAAATTGAAAGTGACAAAATGGTTCAAAAATTGACCTTTACCTATACGGTTACGTATGT
GAATCCTAAAACAAATGATCTTGGTAATATATCAAGTATGCGTCCTGGATATTCTATAATTCAAGTTCAACACAAA

TACATCCTTTACATTTACCCCTTACGCTGCTAGAACAGATAGAATTGGAATTAACTACTTCAATGGTGGAGGAAAGGTAGTTG AATCTAGCACGACCAGTCAGTCACTTTCACAGTCTAAGTCACTCTCAGTAAGTGCTAGTCAAAGCGCCTCAGCTTCAGCATCA ACAAGTGCGTCGGCTTCAGCATCAACCAGTGCCTCGGCTTCAGCGTCAACCAGTGCGTCAGCTTCAGCAAGTACCAGTGCTT AGCATCAACGAGTGCTTCGGCTTCAGCAAGCACAAGCGCCTCGGGTTCAGCATCAACGAGTACGTCAGCTTCAGCGTCAACC AGTGCTTCAGCCTCAGCATCAACAAGTGCGTCAGCCTCAGCAAGTATCTCAGCGTCTGAATCGGCATCAACGAGTGCGTCTG AGTCAGCATCAACGAGTACGTCAGCCTCAGCAAGCACCTCAGCTTCTGAATCGGCCTCAACCAGTGCGTCAGCCTCAGCATC GACAAGCGCCTCAGCTTCAGCAAGTACCAGTGCTTCAGCCTCAGCGTCGACAAGTGCGTCGGCCTCAACCAGTGCATCTGAA TCGGCATCAACCAGTGCGTCAGCCTCAGCAAGTACTAGTGCATCGGCTTCAGCATCAACCAGTGCCTCGGCTTCAGCGTCAA CCAGTGCGTCAGCTTCAGCAAGTACCAGTGCTTCAGTCTCAGCATCAACAAGTGCTTCAGCCTCAGCATCGACAAGTGCCTC GGCTTCAGCAAGCACATCAGCATCTGAATCAGCGTCGACAAGCGCCTCAGCTTCAGCAAGTACCAGTGCGTCAGCTTCAGCA AGCATCAACGAGTACGTCAGCTTCAGCGTCAACCAGTGCTTCAGCCTCAGCATCAACAAGTGCGTCAGCCTCAGCAAGTATC AATCGGCCTCAACCAGTGCGTCAGCCTCAGCATCGACAAGCGCCTCAGCTTCAGCAAGTACCAGTGCTTCAGCCTCAGCTCG ACAAGTGCGTCGGCCTCAACCAGTGCATCTGAATCGGCATCAACCAGTGCGTCAGCCTCAGCAAGTACTAGTGCATCAGCTT CAGCATCAACGAGTGCATCGGCTTCAGCATCAACCAGTGCCTCGGCTTCAGCGTCAACCAGTGCGTCAGCTTCAGCAAGTAC CAGTGCTTCAGTCTCAGCATCAACAAGTGCTTCAGCCTCAGCATCGACAAGTGCcTCGGCTTCAGCAAGCACATCAGCATCTG AATCAGCGTCGACAAGCGC&TCAGCTTCAGCAAGTACCAGTGCGTCAGCCTCAGCGTCGACAAGTGCGTCAGCCTCAGCAAG TACTAGTGCATCAGCTTCAGCATCAACGAGTGCATCGGCTTCGGCGTCAACCAGTGCATCAGAGTCAGCAAGTACCAGTGCG CCTCAACCAGTGCGTCAGCCTCAGCAAGTATCTCAGCGTCTGAATCGGCATCAACGAGTGCGTCCGCTTCAGCAAGTACTAG CGCCTCAGCCTCAGCGTCAACAAGTGCATCGGCTTCAGCGTCAACGAGTGCGTCTGAATCGGCATCAACGAGTGCGTCCGCT TCAGCAAGTACTAGCGCCTCAGCCTCAGCGTCAACAAGTGCATCGGCTTCAGCATCAACGAGTGCGTCCGCTTCAGCAAGTA CTAGCGCCTCAGCCTCAGCGTCAACAAGTGCATCGGCTTCAGCGTCAACGAGTGCGTCTGAGTCAGCATCAACGAGTGCGTC AGCCTCAGCAAGCACATCAGCTTCTGAATCTGCATCAACCAGTGCGTCAGCCTCAGCATCGACAAGCGCCTCAGCTTCAGCA AGTACCAGTGCGTCAGCCTCAGCGTCGACAAGTGCGTCGGCTTCAGCAAGTACCAGTGCGTCAGCCTCAGCAAGTACCAGTG CGTCAGCCTCAGCGTCGACAAGTGCGTCGGCCTCAACCAGTGCATCTGAATCGGCATCAACCAGTGCGTCAGCCTCAGCAAG TACTAGTGCATCAGCTTCAGCATCAACGAGTGCATCGGCTTCAGCATCAACCAGTGCATCAGAGTCAGCAAGTACCAGTGCG TCAGTTCCGCATCAACAAGTGCCTCGGCTTCAGCAAGTACTAG

4183.1 (SEQ. ID. NO. 322)

GCCCTTGGCCGTAGTGCTGGCGTTTTCCGCTATCAAAACCTTTGGAAAACTGGGCAAGAATAACAAGTACAACTTTATCGGTT TTGGGAAATTGTTCCAACTTGGTGGAACGGGTGATTATGCTCAGTTATTTACTTCAATCATTTCAAATCCAGCCATTGCCCTA GGAGCTCAAGCGGCCTTTATCCTATTGAATATCTTCATTGTATCACGTGGGGTTCAAAAAGGGATTGAAAGAGCTTCGAAAG TCATGATGCCCCTGCTCTTTATCGTCTTTGTTTTTATCATCGGTCGCTCTCTCAGTTTGCCAAATGCCATGGAAGGGGTTCTTT GGGTTACAGTCATGTTGACCTATGCTTCTTACTTAGACAAGAAAACCAATCTAGTCCAGTCAGGAATCTCCATCGTAGCCATG AATATCTCGATATCCATCATGGCAGGTCTAGCCATTTTCCAAGCTCGATCCCCCTTCAATATCCAGTCTGAAGGGGGACCCAG CCTGCTCTTTATCGTCTTGCCTCAACTCTTTGACAAGATGCCTTTTGGAACCATTTTCTACGTCCTCTTCCTCTTCCTT AATGGAGTGTTATTTTAGGAATTTTGACCTTTGTCTTTGGCATTCCTTCAGCCCTATCTTACGGTGTCATGGCGGATGTTCACA TTTTTGGTAAGACCTTCTTTGACGCTATGGACTTCTTGGTTTCCAATCTCCTCATGCCATTTGGAGCTCTCTACCTTTCACTTTT TACAGGCTATATCTTTAAAAAGGCTCTTGCAATGGAGGAACTCCATCTCGATGAAAGAGCATGGAAACAAGGACTGTTCCAA GTCTGGCTCTTCCTTCGTTTCGTTTCGTCATTCCAATCATCATCATTGTGGTCTTCATTGCCCAATTTATGTAATCAAA AAGGACTTGAGTAG

4183.5 (SEQ. ID. NO. 323)

4183.6 (SEQ. ID. NO. 324)

4183.7 (SEQ. ID. NO. 325)

4183.8 (SEQ. ID. NO. 326)

4185.3 (SEQ. ID, NO. 327)

4186.1 (SEQ. ID. NO. 328)

4186.2 (SEQ. ID. NO. 329)

4187.2 (SEQ. ID. NO. 330)

4188.1 (SEQ. ID. NO. 331)

4188.2 (SEQ. ID. NO. 332)

4188.5 (SEQ. ID. NO. 333)

4188.10 (SEQ. ID. NO. 334)

ATGTTTAAAGTTTTACAAAAAGTTGGAAAAGCTTTTATGTTACCTATAGCTATACTTCCTGCAGCAGGTCTACTTTTGGGGAT TGAGCTCTGCAGGAGAGGTTGTATTCAGTAATTTGTCACTACTTCTCTGTGTGGGGATTATGTATTGGCTTAGCGAAACGAGAT AAAGGAACCGCTGCGTTAGCAGGAGTAACTGGTTACTTAGTTATGACTGCAACGATCAAAGCTTTGGTAAAACTTTTTATGG CAGAAGGATCTGCAATTGATACTGGAGTTATTGGAGCATTAGTTGTCGGAATAGTTGCCGTATATTTGCACAACCGATATAA CAATATTCAATTACCITCCGCTTTAGGATTCTTTGGAGGTTCACGCTTCGTTCCTATTGTTACATCGTTCTCTTCTATCTTGATT GGCTTTGTCTTCTTTGTTATTTGGCCACCTTTCCAACACTTCTTGTTTCTACAGGTGGATATATTTCTCAGGCGGGTCCAATT GGAACTITTCTATATGGATTITTAATGAGACTTTCTGGAGCAGTAGGCTTACATCATATAATTTACCCTATGTTTTGGTATACT GAACTTGGTGGTGTTGAAACTGTTGCAGGACAAACAGTGGTTGGAGCTCAAAAAATATTTTTTTGCTCAATTAGCCGATTTGGC CCATTCTGGATTATTTACAGAAGGAACAAGGTTTTTTGCAGGTCGTTTCTCAACAATGATGTTCGGTTTACCGGCTGCCTGTTT CGGTATTACAGAACCAATTGAATTTATGTTTCTATTCGTCAGTCCGGTTCTATATGTTGTTCACGCATTCCTTGATGGTGTTAG CTTCTTTATTGCAGACGTCTTAAATATTTCAATAGGAAACACATTTTCAGGAGGTGTAATCGATTTCACTTTATTTGGAATTTT GCAGGGGAACGCTAAGACGAATTGGGTTCTTCAGATTCCATTTTGGACTTATTTTGGAGTGTTTTTTTATATTATTTTTTAGATG GTTCATTACTCAATTCAACGTTCTAACGCCAGGGCGAGAGAAGAAGTAGATTCTAAAGAAATTTCTGAATCCGCAGATTCA ACTTCAAATACTGCAGATTATTTAAAACAGGATAGCCTACAAATTATCAGAGCCTTGGGTGGATCAAATAATATAGAAGATG TAGATGCTTGTGTGACACGTTTACGTGTAGCTGTAAAAGAAGTTAATCAAGTTGATAAAGCACTTTTAAAACAAATTGGTGC AGTTGATGTCTTAGAAGTGAAGGGTGGCATTCAAGCAATCTATGGAGCAAAAGCAATCTTATATAAAAAATAGTATTAATGAA ATTTTAGGTGTAGATGATTAA

4188.11 (SEQ. ID. NO. 335)

ATGAAATTTAGAAAATTAGCTTGTACAGTACTTGCGGGTGCTGCGGTTCTTGGTCTTGCTGCTTGTGGCAATTCTGGCGGAAG TAAAGATGCTGCCAAATCAGGTGGTGACGGTGCCAAAACAGAAATCACTTGGTGGGCATTCCCAGTATTTACCCAAGAAAAA ACTGGTGACGGTGTTGGAACTTATGAAAAATCAATCATCGAAGCGTTTGAAAAAGCAAACCCAGATATAAAAGTGAAATTG GAAACCATCGACTTCAAGTCAGGTCCTGAAAAAATCACAACAGCCATCGAAGCAGGAACAGCTCCAGACGTACTCTTTGATG CACCAGGACGTATCATCCAATACGGTAAAAACGGTAAATTGGCTGAGTTGAATGACCTCTTCACAGATGAATTTGTTAAAGA TGTCAACAATGAAAACATCGTACAAGCAAGTAAAGCTGGAGACAAGGCTTATATGTATCCGATTAGTTCTGCCCCATTCTAC ATGGCAATGAACAAGAAAATGTTAGAAGATGCTGGAGTAGCAAACCTTGTAAAAGAAGGTTGGACAACTGATGATTTTTGAA AAAGTATTGAAAGCACTTAAAGACAAGGGTTACACACCAGGTTCATTGTTCAGTTCTGGTCAAGGGGGAGACCAAGGAACA CGTGCCTTTATCTCTAACCTTTATAGCGGTTCTGTAACAGATGAAAAAGTTAGCAAATATACAACTGATGATCCTAAATTCGT CAAAGGTCTTGAAAAAGCAACTAGCTGGATTAAAGACAATTTGATCAATAATGGTTCACAATTTGACGGTGGGGCAGATATC CAAAACTTTGCCAACGGTCAAACATCTTTACACAATCCTTTGGGCACCAGCTCAAAATGGTATCCAAGCTAAACTTTTAGAAG CAAGTAAGGTAGAAGTAGAAGTACCATTCCCATCAGACGAAGGTAAGCCAGCTCTTGAGTACCTTGTAAACGGGTTTGC AGTATTCAACAATAAAGACGACAAGAAAGTCGCTGCATCTAAGAAATTCATCCAGTTTATCGCAGATGACAAGGAGTGGGG ACCTAAAGACGTAGTTCGTACAGGTGCTTTCCCAGTCCGTACTTCATTTGGAAAACCTTTATGAAGACAAACGCATGGAAACA ATCAGCGGCTGGACTCAATACTACTCACCATACTACAACACTATTGATGGATTTGCTGAAATGAGAACACTTTGGTTCCCAAT AAAAGCTATGAAACAATAG

4188.12 (SEQ. ID. NO. 336)

4191.1 (SEQ. ID. NO. 337)

ATGAAAAAAACTTTTTTCTTACTGGTGTTAGGCTTGTTTTGCCTTCTTCCACTCTCTGTTTTTGCCATTGATTTCAAGATAAACT CTTATCAAGGGGATTTGTATATTCATGCAGACAATACGGCAGAGTTTAGACAGAAGATAGTTTACCAGTTTGAGGAGGACTT TAAGGGCCAAATCGTGGGACTTGGACGTGCTGGTAAGATGCCTAGCGGGTTTGACATTGACCCTCATCCAAAGATTCAGGCC GCGAAAAACGGTGCAGAACTAGCAGATGTGACTAGCGAAGTAACAGAAGAAGCGGATGGTTATACTGTGAGAGTCTATAAT CCAGGTCAGGAGGGCGACATAGTTGAAGTTGACCTCGTCTGGAACTTAAAAAATTTACTTTTCCTTTATGATGATATCGCTGA ATTAAATTGGCAACCTCTGACAGATAGTTCAGAGTCTATTGAAAAGTTTGAATTTCATGTAAGGGGAGACAAGGGGGCTGAA AAACTCTTTTTCCATACAGGGAAACTTTTTAGAGAGGGAACGATTGAAAAGAGTAACCTTGATTATACTATCCGTTTAGACA ATCTTCCGGCTAAGCGTGGAGTTGAGTTGCATGCCTATTGGCCTCGGACCGATTTTGCTAGCGCTAGGGATCAGGGATTGAA AGGGAATCGTTTAGAAGAGTTTAATAAGATAGAAGACTCGATTGTTAGAGAAAAAGATCAGAGTAAACAACTCGTTACTTG AAATATGCCAAAAATCATCGTCTCTATGAACCACCAATGGAATTAGAGCCTATGGTTTTATCAGAAGCAGTCTACTCGACCTC CTTGGAGGAAGTGAGTCCCTTGGTCAAGGGAGCTGGAAAATTCACCTTTGATCAACTTATTCAAGCTACCTTGCTAGATGTGA TAGACCGTGGGAATGTCTCTATCATTTCAGAAGGAGATGCAGTTGGTTTGAGGCTAGTAAAAGAAGATGGTTTGTCAAGCTT GGTATTGAACCAGATGCAAGAAGGAGTGAGAAAACGAGTTTCCTTCTGGGGGCTCCCAGATTATTATCGTCCTTTAACTGGT GGGGAAAAGGCCTTGCAAGTGGGTATGGGTGCCTTGACTATCCTGCCCCTATTTATCGGATTTGGTTTGTTCTTGTACAGTTT AGACGTTCATGGCTATCTTTACCTCCCTTTGCCAATACTTGGTTTTCTAGGGTTAGTTTTTGTCTGTTTTCTATTATTGGAAGCTT CGACTAGATAATCGTGATGGTGTTCTAAATGAAGCGGGAGCTGAGGTCTACTATCTCTGGACCAGTTTTGAAAATATGTTGC GTGAGATTGCACGATTGGATCAGGCTGAACTGGAAAGTATTGTGGTCTGGAATCGCCTCTTGGTCTATGCGACCTTATTTGGC TATGCGGACAAGGTTAGTCATTTGATGAAGGTTCATCAGATTCAAGTGGAAAATCCAGATATCAATCTCTATGTAGCTTATGG CTTCTGGAAGTGGAAGTTCTGGTGGTGGCTTCTCTGGAGGCGGAGGTGGCGGCAGTATCGGTGCCTTTTAA

4191.2 (SEQ. ID. NO. 338)

ATGAAAAAGTAAGAAAGATATTTCAGAAGGCAGTTGCAGGACTGTGCTGTATATCTCAGTTGACAGCTTTTTTCTTCGATAG TTGCTTTAGCAGAAACGCCTGAAACCAGTCCAGCGATAGGAAAAGTAGTGATTAAGGAGACAGGCGAAGGAGGAGCAGCTTC TAGGAGATGCCGTCTTTGAGTTGAAAAACAATACGGATGGCACAACTGTTTCGCAAAGGACAGAGGCGCAAACAGGAGAAG CGATATTTTCAAACATAAAACCTGGGACATACACCTTGACAGAAGCCCAACCTCCAGTTGGTTATAAACCCTCTACTAAACA ATGGACTGTTGAAGTTGAGAAGAATGGTCGGACGACTGTCCAAGGTGAACAGGTAGAAAATCGAGAAGAGGCTCTATCTGA CCAGTATCCACAAACAGGGACTTATCCAGATGTTCAAACACCTTATCAGATTATTAAGGTAGATGGTTCGGAAAAAAACGGA CAGCACAAGGCGTTGAATCCGAATCCATATGAACGTGTGATTCCAGAAGGTACACTTTCAAAGAGAATTTATCAAGTGAATA ATTTGGATGATAACCAATATGGAATCGAATTGACGGTTAGTGGGAAAAACAGTGTATGAACAAAAAGATAAGTCTGTGCCGCT GGATGTCGTTATCTTGCTCGATAACTCAAATAGTATGAGTAACATTCGAAACAAGAATGCTCGACGTGCGGAAAGAGCTGGT GAGGCGACACGTTCTCTTATTGATAAAATTACATCTGATTCAGAAAATAGGGTAGCGCTTGTGACTTATGCTTCCACTATCTT TGATGGGACCGAGTTTACAGTAGAAAAAGGGGTAGCAGATAAAAACGGAAAGCGATTGAATGATTCTCTTTTTTGGAATTAT GATCAGACGAGTTTTACAACCAATACCAAAGATTATAGTTATTTAAAGCTGACTAATGATAAGAATGACATTGTAGAATTAA AAAATAAGGTACCTACCGAGGCAGAAGACCATGATGGAAATAGATTGATGTACCAATTCGGTGCCACTTTTACTCAGAAAGC TTTGATGAAGGCAGATGAGATTTTGACACAACAAGCGAGACAAAATAGTCAAAAAGTCATTTTCCATATTACGGATGGTGTC CCAACTATGTCGTATCCGATTAATTTTAATCATGCTACGTTTGCTCCATCATATCAAAATCAACTAAATGCATTTTTTAGTAAA GGCAAAGTTACCAGATGTTTACAGATAAGACAGTTTATGAAAAAGGTGCTCCTGCAGCTTTCCCAGTTAAACCTGAAAAATA ATTCTGGCTTATCCGTTTAATTCTAATACTGCTAAAATTACCAATCATGGTGACCCTACAAGATGGTACTATAACGGGAATAT TGCTCCTGATGGGTATGATGTCTTTACGGTAGGTATTGGTATTAACGGAGGATCCTGGTACGGATGAAGCAACGGCTACTAGTT TTATGCAAAGTATTTCTAGTAAAACCTGAAAACTATACCAATGTTACTGACACGACAAAAATATTGGAACAGTTGAATCGTTA TTTCCACACCATCGTAACTGAAAAGAAATCAATTGAGAATGGTACGATTACAGATCCGATGGGTGAGTTAATTGATTTGCAA TTGGGCACAGATGGAAGATTTGATCCAGCAGATTACACTTTAACTGCAAACGATGGTAGTCGCTTGGAGAATGGACAAGCTG TAGGTGGTCCACAAAATGATGGTGGTTTTGTTAAAAAAATGCAAAAGTGCTCTATGATACGACTGAGAAAAGGATTCGTGTAAC AGGTCTGTACCTTGGAACGGATGAAAAAGTTACGTTGACCTACAATGTTCGTTTGAATGATGAGTGTTGTAAGCAATAAATTTT ATGATACCAATGGTCGAACAACCTTACATCCTAAGGAAGTAGAACAGAACACAGTGCGCGACTTCCCGATTCCTAAGATTCG TGATGTGCGGAAGTATCCAGAAATCACAATTTCAAAAGAGAAAAAACTTGGTGACATTGAGTTTATTAAGGTCAATAAAAAT GATAAAAAACCACTGAGAGGTGCGGTCTTTAGTCTTCAAAAACAACATCCGGATTATCCAGATATTTATGGAGCTATTGATC AAAATGGCACTTATCAAAATGTGAGAACAGGTGAAGATGGTAAGTTGACCTTTAAAAATCTGTCAGATGGGAAATATCGATT ATTTGAAAATTCTGAACCAGCTGGTTATAAACCCGTTCAAAATAAGCCTATCGTTGCCTTCCAAATAGTAAATGGAGAAGTC AGAGATGTGACTTCAATCGTTCCACAAGATATACCAGCGGGTTACGAGTTTACGAATGATAAGCACTATATTACCAATGAAC CTATTCCTCCAAAGAGAGAATATCCTCGAACTGGTGGTATCGGAATGTTGCCATTCTATCTGATAGGTTGCATGATGATGGGA GGAGTTCTATTATACACACGGAAACATCCGTAA

4191.3 (SEQ. ID. NO. 339)

ATGAATCAATCAACAATTTTTAACAATGCTTGCTGCCTTATTACTGACAGCGAGTAGCCTGTTTTCAGCTGCAACAGTTTT
TGCGGCTGGGACGACAACAACATCTGTTACCGTTCATAAACTATTGGCAACAGATGGGGATATGGATAAAATTGCAAATGAG
TTAGAAACAGGTAACTATGCTGGTAATAAAGTGGGTGTTCTACCTGCAAATGCAAAAGAAATTGCCGGTGTTATGTTCGTTT
GGACAAATACTAAATAATGAAAATTATTGATGAAAATGGCCAAACTCTAGGAGTGAATATTGATCCACAAAACATTTAAACTCTC
AGGGGCAATGCCGGCAACTGCAATGAAAAAATTAACAGAAGCTGAAGGAGCTAAATTTAACACGGCAAATTTACCAGCTGC
TAAGTATAAAATTTTATGAAATTCACAGTTTATCAACTTATGTCCGGTGAAGATGGAGCAACCTTAACAGGTTCTAAAGCAGTTC
CAATTGAAATTGAATTACCATTGAACGATGTTGTGGATGCCATGTGTATCCAAAAAAATACAGAAGCCAAAAATTGA

TAAAGATTTCAAAGGTAAAGCAAATCCAGATACACCACGTGTAGATAAAGATACACCTGTGAACCACCAAGTTGGAGATGTT GTAGAGTACGAAATTGTTACAAAAATTCCAGCACTTGCTAATTATGCAACAGCAAACTGGAGCGATAGAATGACTGAAGGTT TGGCATTCAACAAAGGTACAGTGAAAGTAACTGTTGATGATGTTGCACTTGAAGCAGGTGATTATGCTCTAACAGAAGTAGC AACTGGTTTTGATTTGAAATTAACAGATGCTGGTTTAGCTAAAGTGAATGACCAAAACGCTGAAAAAACTGTGAAAATCACT TATTCGGCAACATTGAATGACAAAGCAATTGTAGAAGTACCAGAATCTAATGATGTAACATTTAACTATGGTAATAATCCAG ATCACGGGAATACTCCAAAGCCGAATAAGCCAAATGAAAACGGCGATTTGACATTGACCAAGACATGGGTTGATGCTACAG GTGCACCAATTCCGGCTGGAGCTGAAGCAACGTTCGATTTGGTTAATGCTCAGACTGGTAAAGTTGTACAAACTGTAACTTTG ACAACAGACAAAAATACAGTTACTGTTAACGGATTGGATAAAAATACAGAATATAAATTCGTTGAACGTAGTATAAAAGGG TATTCAGCAGATTATCAAGAAATCACTACAGCTGGAGAAATTGCTGTCAAGAACTGGAAAGACGAAAATCCAAAACCACTT GATCCAACAGAGCCAAAAGTTGTTACATATGGTAAAAAGTTTGTCAAAGTTAATGATAAAGATAATCGTTTAGCTGGGGCAG AATTTGTAATTGCAAATGCTGATAATGCTGGTCAATATTTAGCACGTAAAGCAGATAAAGTGAGTCAAGAAGAAGAGAAGCAGTT GGTTGTTACAACAAAGGATGCTTTAGATAGAGCAGTTGCTGCTTATAACGCTCTTACTGCACAACAACAACAAACTCAGCAAGAA AAAGAGAAAGTTGACAAAGCTCAAGCTGCTTATAATGCTGCTGTGATTGCTGCCAACAATGCATTTGAATGGGTGGCAGATA AGGACAATGAAAATGTTGTGAAATTAGTTTCTGATGCACAAGGTCGCTTTGAAATTACAGGCCTTCTTGCAGGTACATATTAC GACTGGACAAGGCATTGAGTATACTGCTGGTTCAGGTAAAGATGACGCTACAAAAGTAGTCAACAAAAAATCACTATCCC ACAAACGGGTGGTATTGGTACAATTATCTTTGCTGTAGCGGGGGCTGCGATTATGGGTATTGCAGTGTACGCATATGTTAAA AACAACAAGATGAGGATCAACTTGCTTAA

4191.4 (SEQ. ID. NO. 340)

4191.5 (SEQ. ID. NO. 341)

4191.6 (SEQ. ID. NO. 342)

4192.3 (SEQ. ID. NO. 343)

ATGAAAAAGTATCAACTICTATICAAAATAAGTGCAGTCTTCTCTTACTTATTTTTCGTATTTAGTCTTTCTCAGCTGACGCTT ATCGTCCAAAACTATTGGCAATTTTCTTCTCAGATAGGCAATTTATTCTGGATTCAAAATATCTTGAGTTTACTTTTTATTGGA GTCATGATTGTGGTTCTTGTTAAGACAGGCCATGGTTATCTCTTCCGCATTCCAAGAAAAAAATGGCTTTGGTATTCGATTTT GACAGTATTAGTGCTAGTGTTCCAGATCTCTTTTAACGTTCAGACAGCTAAACATGTTCAGTCAACTGCGGAAGGTTGGGCTG

4193.1 (SEQ. ID. NO. 344)

4193.3 (SEQ. ID. NO. 345)

4194.1 (SEQ. ID. NO. 346)

4194.4 (SEO. ID. NO. 347)

ATGGTTATCGATCCATTTGCTATCAACGAACTAGACTATTACTTAGTTTCACACTTCCACAGTGATCATATCGACCCATACAC AGCTGCAGCAATTCCAATAATCCTAAGTTAGAGCATGTTAAGTTTATCGGTCCTTACCACTGTGGACGAATCTGGGAAGGAT GGGGTGTTCCAAAAGAACGTATCATCGTTGTTAAACCAGGTGACACTATCGAATTAAAAGATATGAAGATTCATGCAGTAGA ATCATTTGACCGTACTTGCTTGGTAACTCTCCCAGTGAACGGTGCTGATGAGACAGGCGGTGAACTTGCTGGCTTGGCTGTTA CAGATGAAGAAATGGCTCAAAAAGGCTGTAACTATATCTTTGAAAACACCAGGTGGAACCATCTATCATGGTGCAGATTCTCA CTTCTCAAACTATTTTGCAAAACATGGTAAAAATCCGGTAGGAAAATCTCACTCTCAGAAAACATCTATCGACCTTCTTCGATAGATTCTGGTACCAAAAGACTATTATCCCAGTTCACTATGATATCTTGGTACAAATTTTGGAAAATCAGTTCAATTTCCAATTCCAAGATTCTACTAATGAGATTTCCAATTTTGGAAAATCCGTTGCAATACGATTTCCAATTTTTGGAAAATTCCACGTTCACTAATTTACCACGTTCACTAATTTCCAAGACTTTTTGGAAAATACCATCATCCACGTGGTTTTGATTACCTCAAGATCATTATCCCACGTGGTTTTTGATTACCACCTTCATCACTTCAATTTTAAAGCTTTTGGAACACTTTAGTAGAACTTTAGTAGAACTTTAGAACACTTAGTAGAATTCCACCTTTAGTAGAATTCCACCTTTAGTAGAACTTTTGGAACACTTTAGTAGAACTTTTGAACAACACTTAATCCACCGTGGTTTTTGATTTTTGAACAAGACTCAACACTTCAATTTAAAGCTTTTGCTATAAA

4196.2 (SEQ. ID. NO. 348)

ATGTTCCTTTCAGGCTGGTTGTCTAGTTTTGCTAATACTTATATCCATGATTTACTGGGGGTTCTTTTCCCAGATAGTCCATTTT
TAAATGCCTTTGAAAGTGCTATTGCGGCTCCTTTGGTAGAAGAACCCTTGAAATTATTGTCACTTGTTTTTGTTTTTGGCTTTGA
TTCCTGTGCGAAAATTAAAATCTTTGTTTTTACTTGGAATTGCTTCCGGTTTGGGATTCCAAATGATTAAGGATATTGGTTATA
TTCGTACGGATTTGCCAGAGGGCTTTGACTTTACTATTTCCCGAATTTTAAGACCGTATCATCAGGAATTGCCTCTCACTGG
TTTTTTCAGGTCTAGCTGTAGTAGGTGTTTACTTGCTTTACAGAGCCTATAAAGGACAGAGGTTGGCAAGAAACAGGGCC
TTATTTTTCTAGGTTTAGCCTTTGGGAACTCACTTCTTGTTTAACTCTCCTTTTTGTGAGAAACAGAGTTTGCTTTACGGA
TTCCAGTGGTTACGGCTATTGCTCTCTATGGTTTTATCATGCTTTATTGCTTTTGTTGAGAAACACAATGAGTTGATGACCTAG

4197.1 (SEQ. ID. NO. 349)

ATGAAGGTGGAACCACGTTGCGACGTCCTTTCGAGGATGTCGCATTTTTTTATTAGGATACTAATTATGGAGTTGCAAGAATT
AGTGGAGCGCAGTTGGGCAATCCGACAAGCTTATCACGAACTGGAAGTTAAGCATCATGATTCCAAGTGGACGGTAGAAGA
AGACCTCTTGGCTTTATCTAATGATATTGGAAATTTCCAACGACTGGTGATGACAAAGCAAGGACGCTACTATGATGAAACA
CCCTACACACACTGGAACAAAAACTTTCAGAAAATATCTGGTGGCTATTAGAACTTTCTCAACGTTTGGATATAGACATTCTGAC
GGAAATGGAAAACTTCCTCTCTGATAAAGAAAAGCAATTGAACGTTAGGACTTGGAAGTAG

4197.4 (SEQ. ID. NO. 350)

4211.2 (SEQ. ID. NO. 351)

4211.3 (SEQ. ID. NO. 352)

4211.4 (SEQ. ID. NO. 353)

4213.1 (SEQ. ID. NO. 354)

4213.2 (SEQ. ID. NO. 355)

4224.1 (SEQ. ID. NO. 356)

4252.1 (SEQ. ID. NO. 357)

4252.2 (SEQ. ID. NO. 358)

4256.2 (SEQ. ID. NO. 359)

4263.1 (SEQ. ID. NO. 360)

ATGATTGTTTCCATTATTTCTCAAGGATTTGTCTGGGCTATTCTAGGTCTGGGAATCTTTATGACATTTAGGATTTTAAACTTT
CCAGATATGACGACAGAAGGTTCCTTCCCTCTTGGGGGAGCTGTTGCTGTCACTTTGATAACCAAAGGCGTGAACCCATTTTT
AGCGACACTTGTTGCTGTAGGAGCAGGTTGTTTGGCTGGAATGGCAGCAGGCCTTCTTTATACAAAAGGGAAGATCCCAACC
TTGCTCTCAGGGATTTTGGTGATGACTTCTTGTCACTCAATCATCATCTTTGATTATGGGACGTGCGAATTTAGGCCTGCTTTGGA



ACCAAGCAAATTCAGGATGTTTTGCCTTTTGATTCGGATTTGAATCAACTCTTGACAGGTCTCATCTTTGTGAGTATTGTTATT
GCTCTCATGCTCTTTTTCTTGGACACTAAACTCGGACAAGCCTATATTGCTACAGGGGATAATCCTGATATTGCTAGAAGTTT
CGGGATTCATACTGGACGCATGGGCTCATGGGCTTGTCTTATCAAATGGTGTGATTGCCCTTGCAGGTGCCCTCATTGCTC
AGCAAGAAGGTTATGCCGATGTGTCTCGAGGGATCGGGGTTATCGTTGTGGGGCTTGCAAGTTTTATTTGGAGAAGTTTA
TTTCAAGAGTTTGAGCTTGGCAGAGCGTTTGGTTACTATCGTTGTAGGTTCTATCGCTTATCAATTTTTAGTGTGGGCAGTTAT
CGCACTTGGCTTTAATACAAGTTACCTTCGTTTATACAGTGCCTTGATTTTTAGCAGTCTGCCTCATGATTCCAACATTTAAGCA
AACAATCTTGAAAGGAGCCAAGTTAAGCAAATGA

4346.1 (SEQ. ID. NO. 361)

4346.2 (SEQ. ID. NO. 362)

4346.3 (SEQ. ID. NO. 363)

TTTGATCTAGTAGTGAATGTATTTGTAAAAGGAGGAGAATTTTCCCTTGATGCTGTACATCGTGTTCTAAAATCTCAGAGGGC ACTTCAGAGTATTATGAACAGTTTTAAGTTAGCATTTTCACTCATTATTACAGTTAATGTCGTAGGTATTCTTTGTGTTCTATT TACAGAGTACTTTGATATTAAAGGTGCTAAAATTTTAAAATTAGGTTATATGACCTCTTTAATTTATGGAGGAGTGGTTTTAG GGTTTATTGGGTATGGTGCAGTCTTATTCATTATGACATTTTCAGGAACTGCTAATCATACATTGTTTTTAACAAATACAATTC GAAGCGTTGACTATCACACTATTGAGGCTGCTCGAAATATGGGAGCAAAACCATTTACTGTTTTCCGAAAAGTAGTGTTACC AACCTTAATTCCAACTCTATTTGCACTTACTATTATGGTTTTTCTTAGTGGTTTATCTGCAGTAGCAGCACCCATGATTGTTGG TGGTAAAGAATTTCAAACTATAAATCCAATGATTATTACATTTGCAGGGGTGGGGGAATTCTCGTGATTTAGCTGCCCTACTTG CAATTATTTTAGGTATTGCAACTACAATTTTGCTTACTATCATGAATAAGATAGAAAAAGGTGGAAATTATATTTCTATCTCT AAGACTAAAGCGCCTCTTAAAAAAACAAAAATTGCGTCTAAGCCTTGGAATATCATTGCTCACATTGTAGCATATGGATTGT TCACAGTTTTCATGCTTCCACTAATTTTTATAGTATTATACTCATTTACAGATCCAGTTGCAATTCAAACAGGTAACTTAACAT TATTATTGCTGCGACACAGCAACACTCTCGCAGTTGTATTTGCTCGTGTTGTCAGAAAACATAAATCTCGTTTTTGATTTCTT ATTTGAATATGGTGCTCTACTTCCTTGGTTACTACCAAGTACACTTTTAGCAGTAAGTTTATTATTTACTTTTAATCAGCCACA ATTTCTTGTCTTGAATCAGATTTTGGTAGGTAGTTTGGTAATTCTACTTATTGCATATAGTTGTAAAAAATCCCATTTTCTTAT AGAATGGTACGTGCTATTTTATTTAGTGTTGATGATGAGATGGAAGATGCAGCAAGAAGTATGGGTGCTTCACCTTTTTATAC TATGATGAAGGTTATCATTCCATTTATTTTACCGGTTGTTCTCTCTGTTATTGCTTTAAACTTTAACTCTTTATTAACTGACTTC GACTTATCTGTATTCCTTTACCATCCCCTAGCTCAACCATTAGGTATTACGATCTGCAGGTGATGAAACAGCAACATC TAATGCACAAGCTCTGGTATTTGTTTATACAATTGTTCTGATGATTATTTCTGGAACGGTATTATACTTCACACAAAGACCGG GGCGTAAAGTAAGGAAATAA



(SEQ. ID. NO. 1)

MEELVTLDCLFIDRTKIEANANKYSFVWKKTTEKFSAKLQEQIQVYFQEEITPLLIKYAMFDKKQKRGYKESAKNLANWHYNDKE DSYTHPDGWYYRFHHTKYQKTQTDFQQEIKVYYADEPESAPQKGLYMNERYQNLKAKECQALLSPQGRQIFAQRKIDVEPVFGQI KASLGYKRCNLRGKRQVRIDMGLVLMANNLLKYSKMKZ

(SEQ. ID. NO. 2)
MGKGHWNRKRVYSIRKFAVGACSVMIGTCAVLLGGNIAGESVVYADETLITHTAEKPKEEKMIVEEKADKALETKNIVERTEQSE PSSTEAIASEKKEDEAVTPKEEKVSAKPEEKAPRIESQASNOEKPLKEDAKAVTNEEVNOMIEDRKVDFNONWYFKLNANSKEAIK PDADVSTWKKLDLPYDWSIFNDFDHESPAQNEGGQLNGGEAWYRKTFKLDEKDLKKNVRLTFDGVYMDSQVYVNGQLVGHYP NGYNQFSYDITKYLQKDGRENVIAVHAVNKQPSSRWYSGSGIYRDVTLQVTDKVHVEKNGTTILTPKLEEQQHGKVETHYTSKIV NTDDKDHELVAEYQIVERGGHAVTGLVRTASRTLKAHESTSLDAILEVERPKLWTVLNDKPALYELITRVYRDGQLVDAKKDLFG YRYYHWTPNEGFSLNGERIKFHGVSLHHDHGALGAEENYKAEYRRLKOMKEMGVNSIRTTHNPASEOTLOIAAELGLLVOFEAF DTWYGGKKPYDYGRFFEKDATHPEARKGEKWSDFDLRTMVERGKNNPAIFMWSIGNEIGEANGDAHSLATVKRLVKVIKDVDK TRYVTMGADKFRFGNGSGGHEKIADELDAVGFNYSEDNYKALRAKHPKWLIYGSETSSATRTRGSYYRPERELKHSNGPERNYE QSDYGNDRVGWGKTATASWTFDRDNAGYAGOFIWTGTDYIGEPTPWHNONOTPVKSSYFGIVDTAGIPKHDFYLYOSOWVSVK KKPMVHLLPHWNWENKELASKVADSEGKIPVRAYSNASSVELFLNGKSLGLKTFNKKQTSDGRTYQEGANANELYLEWKVAYQ PGTLEAIARDESGKEIARDKITTAGKPAAVRLIKEDHAIAADGKDLTYIYYEIVDSQGNVVPTANNLVRFQLHGOGQLVGVDNGEQ ASRERYKAQADGSWIRKAFNGKGVAIVKSTEQAGKFTLTAHSDLLKSNQVTVFTGKKEGQEKTVLGTEVPKVQTIIGEAPEMPTT VPFVYSDGSRAERPVTWSSVDVSKPGIVTVKGMADGREVEARVEVIALKSELPVVKRIAPNTDLNSVDKSVSYVLIDGSVEEYEVD KWEIAEEDKAKLAIPGSRIQATGYLEGQPIHATLVVEEGNPAAPAVPTVTVGGEAVTGLTSQKPMQYRTLAYGAKLPEVTASAKN AAVTVLQASAANGMRASIFIQPKDGGPLQTYAIQFLEEAPKIAHLSLQVEKADSLKEDQTVKLSVRAHYQDGTQAVLPADKVTFS TSGEGEVAIRKGMLELHKPGAVTLNAEYEGAKDQVELTIQANTEKKIAQSIRPVNVVTDLHOEPSLPATVTVEYDKGFPKTHKVT WQAIPKEKLDSYQTFEVLGKVEGIDLEARAKVSVEGIVSVEEVSVTTPIAEAPQLPESVRTYDSNGHVSSAKVAWDAIRPEQYAKE GVFTVNGRLEGTQLTTKLHVRVSAQTEQGANISDQWTGSELPLAFASDSNPSDPVSNVNDKLISYNNQPANRWTNWNRTNPEASV GVLFGDSGILSKRSVDNLSVGFHEDHGVGVPKSYVIEYYVGKTVPTAPKNPSFVGNEDHVFNDSANWKPVTNLKAPAQLKAGEM NHFSFDKVETYAVRIRMVKADNKRGTSITEVQIFAKQVAAAKQGQTRIQVDGKDLANFNPDLTDYYLESVDGKVPAVTASVSNN GLATVVPSVREGEPVRVIAKAENGDILGEYRLHFTKDKSLLSHKPVAAVKQARLLQVGQALELPTKVPVYFTGKDGYETKDLTVE WEEVPAENLTKAGQFTVRGRVLGSNLVAEITVRVTDKLGETLSDNPNYDENSNQAFASATNDIDKNSHDRVDYLNDGDHSENRR WTNWSPTPSSNPEVSAGVIFRENGKIVERTVTQGKVQFFADSGTDAPSKLVLERYVGPEFEVPTYYSNYQAYDADHPFNNPENWE AVPYRADKDIAAGDEINVTFKAIKAKAMRWRMERKADKSGVAMIEMTFLAPSELPOESTOSKILVDGKELADFAENRODYOITYK GQRPKVSVEENNQVASTVVDSGEDSFPVLVRLVSESGKQVKEYRIHLTKEKPVSEKTVAAVQEDLPKIEFVEKDLAYKTVEKKDS TLYLGETRVEQEGKVGKERIFTAINPDGSKEEKLREVVEVPTDRIVLVGTKPVAQEAKKPQVSEKADTKPIDSSEASQTNKAQLPST GSAASQAAVAAGLTLLGLSAGLVVTKGKKEDZ

(SEQ. ID. NO. 3) MKIMKKKYWTLAILFFCLFNNSVTAQEIPKNLDGNITHTQTSESFSESDEKQVDYSNKNQEEVDQNKFRIQIDKTELFVTTDKHLEK NCCKLELEPQINNDIVNSESNNLLGEDNLDNKIKENVSHLDNRGGNIEHDKDNLESSIVRKYEWDIDKVTGGGESYKLYSKSNSKV SIAILDSGVDLONTGLLKNLSNHSKNYVPNKGYLGKEEGEEGIISDIODRLGHGTAVVAOIVGDDNINGVNPHVNINVYRIFGKSSA SPDWIVKAIFDAVDDGNDIINLSTGQYLMIDGEYEDGTNDFETFLKYKKAIDYANQKGVIIVAALGNDSLNVSNQSDLLKLISSRKK VRKPGLVVDVPSYFSSTISVGGIDRLGNLSDFSNKGDSDAIYAPAGSTLSLSELGLNNFINAEKYKEDWIFSATLGGYTYLYGNSFA **APKVSGAIAMIIDKYKLKDQPYNYMFVKKFWKKHYQZ**

(SEQ. ID. NO. 4)

MKKTWKVFLTLVTALVAVVLVACGQGTASKDNKEAELKKVDFILDWTPNTNHTGLYVAKEKGYFKEAGVDVDLKLPPEESSSD LVINGKAPFAVYFQDYMAKKLEKGAGITAVAAIVEHNTSGIISRKSDNVSSPKDLVGKKYGTWNDPTELAMLKTLVESQGGDFEK VEKVPNNDSNSITPIANGVFDTAWIYYGWDGILAKSOGVDANFMYLKDYVKEFDYYSPVIIANNDYLKDNKEEARKVIOAIKKGY QYAMEHPEEAADILIKNAPELKEKRDFVIESQKYLSKEYASDKEKWGQFDAARWNAFYKWDKENGILKEDLTDKGFTNEFVKZ

MKRTWRNSFVTNLNTPFMIGNIEIPNRTVLAPMAGVTNSAFRTIAKELGAGLVVMEMVSDKGIOYNNEKTLHMLHIDEGENPVSI QLFGSDEDSLARAAEFIQENTKTDIVDINMGCPVNKIVKNEAGAMWLKDPDKIYSIINKVQSVLDIPLTVKMRTGWADPSLAVENA LAAEAAGVSALAMHGRTREOMYTGHADLETLYKVAOALTKIPFIANGDIRTVOEAKORIEEVGADAVMIGRAAMGNPYLFNOIN HYFETGEILPDLTFEDKMKIAYEHLKRLINLKGENVAVREFRGLAPHYLRGTSGAAKLRGAISQASTLAEIETLLQLEKAZ

(SEQ. ID. NO. 6)

MIKNPKLLTKSFLRSFAILGGVGLVIHIAIYLTFPFYYIQLEGEKFNESARVFTEYLKTKTSDEIPSLLQSYSKSLTISAHLKRDIVDKR LPLVHDLDIKDGKLSNYIVMLDMSVSTADGKQVTVQFVHGVDVYKEAKNILLLYLPYTFLVTIAFSFVFSYFYTKRLLNPLFYISEV TSKMQDLDDNIRFDESRKDEVGEVGKQINGMYEHLLKVIYELESRNEQIVKLONQKVSFVRGASHELKTPLASLRIILENMOHNIG DYKDHPKYIAKSINKIDQMSHLLEEVLESSKFQEWTECRETTVKPVLVDILSRYQELAHSIGVTIENQLTDATRVVMSLRALDKVLT NLISNAIKYSDKNGRVIIŠEQDGYLSIKNTCAPLSDQELEHLFDIFYHSQIVTDKDESSGLGLYIVNNILESYQMDYSFLPYEHGMEFK ISLZ

(SEQ. ID. NO. 7)

MYLGDLMEKAECGQFSILSFLLQESQTTVKAVMEETGFSKATLTKYVTLLNDKALDSGLELAIHSEDENLRLSIGAATKGRDIRSLF LESAVKYOILVYLLYHOOFLAHOLAOELVISEATLGRHLAGLNOILSEFDLSIONGRWRGPEHOIHYFYFCLFRKVWSSOEWEGHM QKPERKQEIANLEEICGASLSAGQKLDLVLWAHISQQRLRVNACQFQVIEEKMRGYFDNIFYLRLLRKVPSFFAGQHIPLGVEDGE MMIFFSFLLSHRILPLHTMEYILGFGGQLADLLTQLIQEMKKEELLGDYTEDHVTYELSQLCAQVYLYKGYILQDRYKYQLENRHP YLLMEHDFKETAEEIFHALPAFOOGTDLDKKILWEWLOLIEYMAENGGOHMRIGLDLTSGFLVFSRMAAILKRYLEYNRFITIEAY DPSRHYDLLVTNNPIHKKEQTPVYYLKNDLDMEDLVAIRQLLFTZ

MEFSKKTRELSIKKMOERTLDLLIIGGGITGAGVALOAAASGLETGLIEMODFAEGTSSRSTKLVHGGLRYLKOFDVEVVSDTVSE RAVVQQIAPHIPKSDPMLLPVYDEDGATFSLFRLKVAMDLYDLLAGVSNTPAANKVLSKDQVLEROPNLKKEGLVGGGVYLDFR NNDARLVIENIKRANQDGALIANHVKAEGFLFDESGKITGVVARDLLTDQVFEIKARLVINTTGPWSDKVRNLSNKGTQFSQMRPT KGVHLVVDSSKIKVSQPVYFDTGLGDGRMVFVLPRENKTYFGTTDTDYTGDLEHPKVTQEDVDYLLGIVNNRFPESNITIDDIESS WAGLRPLIAGNSASDYNGGNNGTISDESFDNLIATVESYLSKEKTREDVESAVSKLESSTSEKHLDPSAVSRGSSLDRDDNGLLTLA GGKITDYRKMAEGAMERVVDILKAEFDRSFKLINSKTYPVSGGELNPANVDSEIEAFAQLGVSRGLDSKEAHYLANLYGSNAPKV FALAHSLEQAPGLSLADTLSLHYAMRNELTLSPVDFLLRRTNHMLFMRDSLDSIVEPILDEMGRFYDWTEEEKATYRADVEAALA NNDLAELKNZ

(SEQ. ID. NO. 9)

MMNELFGEFLGTLILILLGNGVVAGVVLPKTKSNSSGWIVITMGWGIAVAVAVFVSGKLSPAYLNPAVTIGVALKGGLPWASVLP YILAQFAGAMLGQILVWLQFKPHYEAEENAGNILATFSTGPAIKDTVSNLISEILGTFVLVLTIFALGLYDFQAGIGTFAVGTLIVGIG LSLGGTTGYALNPARDLGPRIMHSILPIPNKGDGDWSYAWIPVVGPVIGAALAVLVFSLFZ

(SEO. ID. NO. 10)

MTKKKIERISVIHREKILWLKWYFMRDKEQPKYSVLERKMFDAAKNQDMLAYQKYATIKQITDIRVQTSEADILEAVKEVYVYNH MNVIGACQRILFISQSPAYDKLNKWFNIYSDLYFSVVPLPKMGVYHEMVGIZ

(SEQ. ID. NO. 11)

MKNSNEAEMKLLYTDIRTSLTEILTREAEELVAAGKRVFYIAPNSLSFEKERAVLEYLSQQASFSITVTRFAQMARYLVLNDLPAKT TLDDIGLGLAFYKCLAELDPKDLRVYGAIKQDPQLIQQLIELYHEMTKSQMSFLDLENLTDEDKRADLLLIFEKVTAYLNQGQLAQ ESQLSHLIEAIENDKVSSDFNQIALVIDGFTRFSAEEERVVDLLHGKGVEIVIGAYASKKAYTSPFSEGNLYQASVKFLHHLASKYQT PAQDCSQTHEKMDSFDKASRLLESSYDFSELALDVDEKDRENLQIWSCLTQKEELELVARSIRQKLHENSDLSYKHFRILLGDVAS YQLSLKTIFDQYQIPFYLGRSEAMAHPLTQFVESILALKRYRFRQEDLINLLRTDLYTDLSQSDIDAFEQYIRYLGINGLPAFQQTFT KSHHGKFNLERLNVLRLRILAPLETLFASRKQKAEKLLQKWSVFLKEGAVTKQLQDLTTTLEAVEQERQAEVWKAFCHVLEQFAT VFAGSQVSLEDFLALLHSGMSLSQYRTIPATVDTVLVQSYDLIAPLTADFVYAIGLTQDNLPKISQNTSLLTDEERQNLNQATEEGV QLLIASSENLKKNRYTMLSLVNSARKQLFLSAPSLFNESESKESAYLQELIHFGFRRREKRMNHKGLSKEDMGSYHSLLSSLVAYH QQGEMSDTEQDLTFVKVLSRVIGKKLDQQGLENPAIPTSPSSKTLAKDTLQALYPAKQEFYLSTSGLTEFYRNEYSYFLRYVLGLQ EELRLHPDARSHGNFLHRIFERALQLPNEDSFDQRLEQAIQETSQEREFEAIYQESLEAQFTKEVLLDVARTTGHILRHNPAIETIKEE ANFGGKDQAFIQLDNGRSVFVRGKVDRIDRLKANGAIGVVDYKSSLTQFQFPHFFNGLNSQLPTYLAALKREGEQNFFGAMYLEM AEPVQSLMAVKSLAGAVVEASKSMKYQGLFLEKESSYLGEFYNKNKANQLTDEEFQLLLDYNAYLYKKAAEKILAGRFAINPYT ENGRSIAPYVQQHQAITGFEANYHLGQARFLEKLDLADGKRLVGEKLKQAWLEKIREELNRZ

(SEQ. ID. NO. 12)

MKLIPFLSEEEIQKLQEAEANSSKEQKKTAEQIEAIYTSAQNILVSASAGSGKTFVMAERILDQLARGVEISQLFISTFTVKAATELKE RLEKKISKKIQETDDVDLKQHLGRQLADLPNAAIGTMDSFTQKFLGKHGYLLDIAPNFRILQNQSEQLILENEVFHEVFEAHYQGK QKETFSHLLKNFAGRGKDERGLRQQVYKIYDFLQSTSNPQKWLSESFLKGFEKADFTSEKEKLTEQIKQALWDLESFFRYHLDND AKEFAKAAYLENVQLILDEIGSLNQESDSQAYQAVLARVVAISKEKNGRALTNASRKADLKPLADAYNEERKTQFAKLGQLSDQI AILDYQERYHGDTWKLAKTFQSFMSDFVEAYRQRKRQENAFEFADISHYTIEILENFPQVRESYQERFHEVMVDEYQDTNHIQER MLELLSNGHNRFMVGDIKQSIYRFRQADPQIFNEKFQRYAQNPQEGRLIILKENFRSSSEVLSATNDVFERLMDQEVGEINYDNKH QLVFANTKLTPNPDNKAAFLLYDKDDTGEEEESQTETKLTGEMRLVIKEILKLHQEKGVAFKEIALLTSSRSRNDQILLALSEYGIP VKTDGEQNNYLQSLEVQVMLDTLRVIHNPLQDYALVALMKSPMFGFDEDELARLSLQKAEDKVHENLYEKLVNAQKMASSQKG LIHTALAEKLKQFMDILASWRLYAKTHSLYDLIWKIYNDRFYYDYVGALPNGPARQANLYALALRADQFEKSNFKGLSRFIRMID QVLEAQHDLASVAVAPPKDAVELMTIHKSKGLEFPYVFILNMDQDFNKQDSMSEVILSRQNGLGVKYIAKMETGAVEDHYPKTIK LSIPSLTYRQNEEELQLASYSEQMRLLYVAMTRAEKKLYLVGKGSREKLESKEYPAAKNGKLNSNTRLQARNFQDWLWAISKVFT KDKLNFSYRFIGEDQLTREAIGELETKSPLQDSSQADNRQSDTIKEALEMLKEVEVYNTLHRAAIELPSVQTPSQIKKFYEPVMDME GVEIAGQGQSVGKKISFDLPDFSTKEKVTGAEIGSATHELMQRIDLSQQLTLASLTETLKQVQTSQAVRDKINLDKILAFFDTVLGQ EILANTDHLYREQPFSMLKRDQKSQEDFVVRGILDGYLLYENKIVLFDYKTDRYDEPSQLVDRYRGQLALYEEALSRAYSIENIEK YLLLGKDEVQVVKVZ

(SEQ. ID. NO. 13)

MELARHAESLGVDAIATIPPIYFRLPEYSVAKYWNDISSAAPNTDYVIYNIPQLAGVALTPSLYTEMLKNPRVIGVKNSSMPVQDIQ TFVSLGGEDHIVFNGPDEQFLGGRLMGARAGIGGTYGAMPELFLKLNQLIADKDLETARELQYAINAIIGKLTSAHGNMYGVIKEV LKINEGLNIGSVRSPLTPVTEEDRPVVEAAAALIRETKERFLZ

(SEQ. ID. NO. 14)

MYKTKCLREKLVLFLKIFFPILIYQFANYSASFVDTAMTGQYNTMDLAGVSMATSIWNPFFTFLTGIVSALVPIIGHHLGRGKKEEV ASDFYQFIYLALGLSVVLLGMVLFLAPIILNHIGLEAAVAAVAVYLWFLSIGIIPLLLFSVIRSLLDSLGLTKLSMYLMLLLLPLNSG FNYLLIYGAFGVPELGGAGAGLGTSLAYWVLLGISVLVLFKQEKLKALHLEKRIPLNMDKIKEGVRLGLPIGGTVFAEVAIFSVVGL IMAKFSPLIIASHQSAMNFSSLMYAFPMSISSAMAIVVSYEVGAKRFDDAKTYIGLGRWTALIFAAFTLTFLYIFRGNVASLYGNDP KFIDLTVRFLTYSLFFQLADTFAAPLQGILRGYKDTVIPFYLGLLGYWGVAIPVYAIZ

<u>(SEQ. ID. NO. 15)</u>

MSTLAKIEALLFVAGEDGIRVRQLAELLSLPPTGIQQSLGKLAQKYEKDPDSSLALIETSGAYRLVTKPQFAEILKEYSKAPINQSLS RAALETLSIIAYKQPITRIEIDAIRGVNSSGALAKLQAFDLIKEDGKKEVLGRPNLYVTTDYFLDYMGINHLEELPVIDELEIQAQESQ LFGERIEEDENOZ

(SEQ. ID. NO. 16)

MDTMISRFFRHLFEALKSLKRNGWMTVAAVSSVMITLTLVAIFASVIFNTAKLATDIENNVRVVYYIRKDVEDNSQTIEKEGQTVT NNDYHKVYDSLKNMSTVKSVTFSSKEEQYEKLTEIMGDNWKIFEGDANPLYDAYIVEANTPNDVKTIAEDAKKIEGVSEVQDGG ANTERLFKLASFIRVWGLGIAALLIFIAVFLISNTIRITIISRSREIQIMRLVGAKNSYIRGPFLLEGAFIGLLGAIAPSVLVFIVYQIVYQS VNKSLVGQNLSMISPDLFSPLMIALLFVIGVFIGSLGSGISMRRFLKIZ

(SEQ. ID. NO. 17)

MKKVRFIFLALLFFLASPEGAMASDGTWQGKQYLKEDGSQAANEWVFDTHYQSWFYIKADANYAENEWLKQGDDYFYLKSGG YMAKSEWVEDKGAFYYLDQDGKMKRNAWVGTSYVGATGAKVIEDWVYDSQYDAWFYIKADGQHAEKEWLQIKGKDYYFKS GGYLLTSQWINQAYVNASGAKVQQGWLFDKQYQSWFYIKENGNYADKEWIFENGHYYYLKSGGYMAANEWIWDKESWFYLK FDGKMAEKEWVYDSHSQAWYYFKSGGYMTANEWIWDKESWFYLKSDGKIAEKEWVYDSHSQAWYYFKSGGYMTANEWIWD KESWFYLKSDGKIAEKEWVYDSHSQAWYYFKSGGYMAKNETVDGYQLGSDGKWLGGKTTNENAAYYQVVPVTANVYDSDGE KLSYISQGSVVWLDKDRKSDDKRLAITISGLSGYMKTEDLQALDASKDFIPYYESDGHRFYHYVAQNASIPVASHLSDMEVGKKY YSADGLHFDGFKLENPFLFKDLTEATNYSAEELDKVFSLLNINNSLLENKGATFKEAEEHYHINALYLLAHSALESNWGRSKIAKD KNNFFGITAYDTTPYLSAKTFDDVDKGILGATKWIKENYIDRGRTFLGNKASGMNVEYASDPYWGEKIASVMMKINEKLGGKDZ

(SEQ. ID. NO. 18)

MKKVLQKYWAWAFVVIPLLLQAIFFYVPMFQGAFYSFTNWTGLTYNYKFVGLNNFKLLFMDPKFMNAIGFTAIIAIAMVVGEIAL GIFIARVLNSKIKGQTFFRAWFFFPAVLSGLTVALIFKQVFNYGLPAIGNALHIEFFQTSLLGTKWGAIFAAVFVLLWQGVAMPIIIFL AGLQSIPTEITEAARIDGATSKQVFWNIELPYLLPSVSMVFILALKGGLTAFDQVFAMTGGGPNNATTSLGLLVYNYAFKNNQFGY ANAIAVILFFLIVVISIIQLRVSKKFEIZ

(SEQ. ID. NO. 19)

MMKQDERKALIGKYILLILGSVLILVPLLATLFSSFKPTKDIVDNFFGFPTNFTWDNFSRLLADGIGGYYWNSVVITVLSLLAVMIFIP MAAYSIARNMSKRKAFTIMYTLLILGIFVPFQVIMIPITVMMSKLGLANTFGLILLYLTYAIPQTLFLYVGYIKISIPESLDEAAEIDGA NQFTTYFRIIFPMMKPMHATTMIINALWFWNDFMLPLLVLNRDSKMWTLPLFQYNYAGQYFNDYGPSFASYVVGIISITIVYLFFQ RHIISGMSNGAVKZ

(SEQ. ID. NO. 20)

MKSILQKMGEHPMLLLFLSYSTVISILAQNWMGLVASVGMFLFTIFFLHYQSILSHKFFRLILQFVLFGSVLSAAFASLEHFQIVKKF NYAFLSPNMQVWHQNRAEVTFFNPNYYGIICCFCIMIAFYLFTTTKLNWLKVFCVIAGFVNLFGLNFTQNRTAFPAIIAGAIIYLFTTI KNWKAFWLSIGVFAIGLSFLFSSDLGVRMGTLDSSMEERISIWDAGMALFKQNPFWGEGPLTYMNSYPRIHAPYHEHAHSLYIDTI LSYGIVGTILLVLSSVAPVRLMMDMSQESGKRPIIGLYLSFLTVVAVHGIFDLALFWIQSGFIFLLVMCSIPLEHRMLVSDMTDZ

(SEQ. ID. NO. 21)

MSKMDVQKIIAPMMKFVNMRGIIALKDGMLAILPLTVVGSLFLIMGQLPFEGLNKSIASVFGANWTEPFMQVYSGTFAIMGLISCFS IAYSYAKNSGVEALPAGVLSVSAFFILLRSSYIPKQGEAIGDAISKVWFGGQGIIGAIIIGLVVGSIYTFFIKRKIVIKMPEQVPQAIAKQ FEAMIPAFVIFLSSMIVYILAKSLTNGGTFIEMIYSAIQVPLQGLTGSLYGAIGIAFFISFLWWFGVHGQSVVNGVVTALLLSNLDAN KAMLASANLSLENGAHIVTQQFLDSFLILSGSGITFGLVVAMLFAAKSKQYQALGKVAAFPAIFNVNEPVVFGFPIVMNPVMFVPFI LVPVLAAVIVYGAIATGFMQPFSGVTLPWSTPAILSGFLVGGWQGVITOLVILAMSTLVYFPFFKVODRLAYQNEIKOSZ

(SEQ. ID. NO. 22)

MKKKDLVDQLVSEIETGKVRTLGIYGHGASGKSTFAQELYQALDSTTVNLLETDPYITSGRHLVVPKDAPNQKVTASLPVAHELES LQRDILACRRVWMSZ

(SEO. ID. NO. 23)

MKKRYLVLTALLALSLAACSQEKTKNEDGETKTEQTAKADGTVGSKSQGAAQKKAEVVNKGDYYSIQGKYDEIIVANKHYPLSK DYNPGENPTAKAELVKLIKAMQEAGFPISDHYSGFRSYETQTKLYQDYVNQDGKAAADRYSARPGYSEHQTGLAFDVIGTDGDL VTEEKAAQWLLDHAADYGFVVRYLKGKEKETGYMAEEWHLRYVGKEAKEIAASGLSLEEYYGFEGGDYVDZ

(SEQ. ID. NO. 24)

MREPDFLNHFLKKGYFKKHAKAVLALSGGLDSMFLFKVLSTYQKELEIELILAHVNHKQRIESDWEEKELRKLAAEAELPIYISNFS GEFSEARARNFRYDFFQEVMKKTGATALVTAHHADDQVETIFMRLIRGTRLRYLSGIKEKQVVGEIEIIRPFLHFQKKDFPSIFHFED TSNQENHYFRNRIRNSYLPELEKENPRFRDAILGIGNEILDYDLAIAELSNNINVEDLQQLFSYSESTQRVLLQTYLNRFPDLNLTKA QFAEVQQILKSKSQYRHPIKNGYELIKEYQQFQICKISPQADEKEDELVLHYQNQVAYQGYLFSFGLPLEGELIQQIPVSRETSIHIRH RKTGDVLIKNGHRKKLRRLFIDLKIPMEKRNSALIIEQFGEIVSILGIATNNLSKKTKNDIMNTVLYIEKIDRZ

(SEQ. ID. NO. 25)

MRKFLIILLPSFLTISKVVSTEKEVVYTSKEIYYLSQSDFGIYFREKLSSPMVYGEVPVYANEDLVVESGKLTPKTSFQITEWRLNK QGIPVFKLSNHQFIAADKRFLYDQSEVTPTIKKVWLESDFKLYNSPYDLKEVKSSLSAYSQVSIDKTMFVEGREFLHIDQAGWVAK ESTSEEDNRMSKVQEMLSEKYQKDSFSIYVKQLTTGKEAGINQDEKMYAASVLKLSYLYYTQEKINEGLYQLDTTVKYVSAVNDF PGSYKPEGSGSLPKKEDNKEYSLKDLITKVSKESDNVAHNLLGYYISNQSDATFKSKMSAIMGDDWDPKEKLISSKMAGKFMEAI YNQNGFVLESLTKTDFDSQRIAKGVSVKVAHKIGDADEFKHDTGVVYADSPFILSIFTKNSDYDTISKIAKDVYEVLKZ

(SEQ. ID. NO. 26)

MKKQNNGLIKNPFLWLLFIFFLVTGFQYFYSGNNSGGSQQINYTELVQEITDGNVKELTYQPNGSVIEVSGVYKNPKTSKEETGIQF FTPSVTKVEKFTSTILPADTTVSELQKLATDHKAEVTVKHESSSGIWINLLVSIVPFGILFFFLFSMMGNMGGGNGRNPMSFGRSKA KAANKEDIKVRFSDVAGAEEEKQELVEVVEFLKDPKRFTKLGARIPAGVLLEGPPGTGKTLLAKAVAGEAGVPFFSISGSDFVEMF VGVGASRVRSLFEDAKKAAPAIIFIDEIDAVGRQRGVGLGGGNDEREQTLNQLLIEMDGFEGNEGIIVIAATNRSDVLDPALLRPGR FDRKVLVGRPDVKGREAILKVHAKNPLAEDVDLKLVAQQTPGFVGADLENVLNEAALVAARRNKSIIDASDIDEAEDRVIAGPS KKDKTVSQKERELVAYHEAGHTIVGLVLSNARVVHKVTIVPRGRAGGYMIALPKEDQMLLSKEDMKEQLAGLMGGRVAEEIIFN VQTTGASNDFEQATQMARAMVTEYGMSEKLGPVQYEGNHAMLGAQSPQKSISEQTAYEIDEEVRSLLNEARNKAAEIIQSNRETH KLIAEALLKYETLDSTQIKALYETGKMPEAVEEESHALSYDEVKSKMNDEKZ

(SEQ. ID. NO. 27)

MKRSSLLVRMVISIFLVFLILLALVGTFYYQSSSSAIEATIEGNSQTTISQTSHFIQSYIKKLETTSTGLTQQTDVLAYAENPSQDKVEG IRDLFLTILKSDKDLKTVVLVTKSGQVISTDDSVQMKTSSDMMAEDWYQKAIHQGAMPVLTPARKSDSQWVISVTQELVDAKGA NLGVLRLDISYETLEAYLNQLQLGQQGFAFIINENHEFVYHPQHTVYSSSSKMEAMKPYIDTGQGYTPGHKSYVSQEKIAGTDWTV LGVSSLEKLDQVRSQLLWTLLGASVTSLLVCLCLVWFSLKRWIAPLKDLRETMLEIASGAQNLRAKEVGAYELREVTRQFNAMLD QIDQLMVAIRSQEETTRQYQLQALSSQINPHFLYNTLDTIIWMAEFHDSQRVVQVTKSLATYFRLALNQGKDLICLSDEINHVRQYL

FIQKQRYGDKLEYEINENVAFDNLVLPKLVLQPLVENALYHGIKEKEGQGHIKLSVQKQDSGLVIRIEDDGVGFQDAGDSSQSQLKRGGVGLONVDORLKLHFGANYHMKIDSRPQKGTKVEIYINRIETSZ

(SEO, ID, NO, 28)

MKRSSLLVRMVISIFLVFLILLALVGTFYYQSSSSAIEATIEGNSQTTISQTSHFIQSYIKKLETTSTGLTQQTDVLAYAENPSQDKVEG IRDLFLTILKSDKDLKTVVLVTKSGQVISTDDSVQMKTSSDMMAEDWYQKAIHQGAMPVLTPARKSDSQWVISVTQELVDAKGA NLGVLRLDISYETLEAYLNQLQLGQQGFAFIINENHEFVYHPQHTVYSSSSKMEAMKPYIDTGQGYTPGHKSYVSQEKIAGTDWTV LGVSSLEKLDQVRSQLLWTLLGASVTSLLVCLCLVWFSLKRWIAPLKDLRETMLEIASGAQNLRAKEVGAYELREVTRQFNAMLD QIDQLMVAIRSQEETTRQYQLQALSSQINPHFLYNTLDTIIWMAEFHDSQRVVQVTKSLATYFRLALNQGKDLICLSDEINHVRQYL FIQKQRYGDKLEYEINENVAFDNLVLPKLVLQPLVENALYHGIKEKEGQGHIKLSVQKQDSGLVIRIEDDGVGFQDAGDSSQSQLK RGGVGLQNVDQRLKLHFGANYHMKIDSRPQKGTKVEIYINRIETSZ

(SEQ. ID. NO. 29)

MFFKLLREALKVKQVRSKILFTIFIVLVFRIGTSITVPGVNANSLNALSGLSFLNMLSLVSGNALKNFSIFALGVSPYITASIVVQLLQ MDILPKFVEWGKQGEVGRRKLNQATRYIALVLAFVQSIGITAGFNTLAGAQLIKTALTPQVFLTIGIILTAGSMIVTWLGEQITDKG YGNGVSMIIFAGIVSSIPEMIQGIYVDYFVNVPSSRITSSIIFVIILIITVLLIIYFTTYVQQAEYKIPIQYTKVAQGAPSSSYLPLKVNPAG VIPVIFASSITAAPAAILQFLSATGHDWAWVRVAQEMLATTSPTGIAMYALLIILFTFFYTFVQINPEKAAETYKRVVPISMEFVLVK VQKNICLNFFVVLQLLVPSSLVZ

(SEQ. ID. NO. 30)

MDIRQVTETIAMIEEQNFDIRTITMGISLLDCIDPDINRAAEKIYQKITTKAANLVAVGDEIAAELGIPIVNKRVSVTPISLIGAATDAT DYVVLAKALDKAAKEIGVDFIGGFSALVQKGYQKGDEILINSIPRALAETDKVCSSVNIGSTKSGINMTAVADMGRIIKETANLSD MGVAKLVVFANAVEDNPFMAGAFHGVGEADVIINVGVSGPGVVKRALEKVRGQSFDVVAETVKKTAFKITRIGQLVGQMASERL GVEFGIVDLSLAPTPAVGDSVARVLEEMGLETVGTHGTTAALALLNDQVKKGGVMACNQVGGLSGAFIPVSEDEGMIAAVQNGS LNLEKLEAMTAICSVGLDMIAIPEDTPAETIAAMIADEAAIGVINMKTTAVRIIPKGKEGDMIEFGGLLGTAPVMKVNGASSVDFISR GGQIPAPIHSFKNZ

(SEQ. ID. NO. 31)

MTQIIDGKALAAKLQGQLAEKTAKLKEETGLVPGLVVILVGDNPASQVYVRNKERSALAAGFRSEVVRVPETITQEELLDLIAKYN QDPAWHGILVQLPLPKHIDEEAVLLAIDPEKDVDGFHPLNMGRLWSGHPVMIPSTPAGIMEMFHEYGIDLEGKNAVVIGRSNIVGK PMAQLLLAKNATVTLTHSRTHNLSKVAAKADILVVAIGRAKFVTADFVKPGAVVIDVGMNRDENGKLCGDVDYEAVAPLASHIT PVPGGVGPMTITMLMEQTYQAALRTLDRKZ

(SEO. ID. NO. 32)

MSKFNRIHLVVLDSVGIGAAPDANNFVNAGVPDGASDTLGHISKTVGLNVPNMAKIGLGNIPRETPLKTVAAESNPTGYATKLEEV SLGKDTMTGHWEIMGLNITEPFDTFWNGFPEEILTKIEEFSGRKVIREANKPYSGTAVIYDFGPRQMETGELIIYTSADPVLQIAAHE DIIPLDELYRICEYARSITLERPALLGRIIARPYVGEPGNFTRTANRRDLAVSPFFPTVLDKLNEAGIDTYAVGKINDIFNGAGINHDM GHNKSNSHGIDTLLKTMGLAEFEKGFSFTNLVDFDALYGHRRNAHGYRDCLHEFDERLPEIIAAMRENDLLLITADHGNDPTYAG TDHTREYIPLLAYSPAFKGNGLIPVGHFADISATVADNFGVETAMIGESFLDKLVZ

(SEQ. ID. NO. 33)

MFISISAGIVTFLLTLVEIPAFIQFYRKAQITGQQMHEDVKQHQAKAGTPTMGGLVFLITSVLVAFFFALFSSQFSNNVGMILFILVLY GLVGFLDDFLKVFRKINEGLNPKQKLALQLLGGVIFYLFYERGGDILSVFGYPVHLGFFYIFFALFWLVGFSNAVNLTDGVDGLASI SVVISLSAYGVIAYVQGQMDILLVILAMIGGLLGFFIFNHKPAKVFMGDVGSLALGGMLAAISMALHQEWTLLIIGIVYVFETTSVM MQVSYFKLTGGKRIFRMTPVHHHFELGGLSGKGNPWSEWKVDFFFWGVGLLASLLTLAILYLMZ

(SEQ. ID. NO. 34)

LFKKNKDILNIALPAMGENFLQMLMGMVDSYLVAHLGLIAISGVSVAGNIITIYQAIFIALGAAISSVISKSIGQKDQSKLAYHVTEA LKITLLLSFLLGFLSIFAGKEMIGLLGTERDVAESGGLYLSLVGGSIVLLGLMTSLGALIRATHNPRLPLYVSFLSNALNILFSSLAIFV LDMGIAGVAWGTIVSRLVGLVILWSQLKLPYGKPTFGLDKELLTLALPAAGERLMMRAGDVVIIALVVSFGTEAVAGNAIGEVLT QFNYMPAFGVATATVMLLARAVGEDDWKRVASLSKQTFWLSLFLMLPLSFSIYVLGVPLTHLYTTDSLAVEASVLVTLFSLLGTP MTTGTVIYTAVWQGLGNARLPFYATSIGMWCIRIGTGYLMGIVLGWGLPGIWAGSLLDNGFRWLFLRYRYQRYMSLKGZ

(SEQ. ID. NO. 35)

MQTQEKHSQAAVLGLQHLLAMYSGSILVPIMIATALGYSAEQLTYLISTDIFMCGVATFLQLQLNKYFGIGLPVVLGVAFQSVAPLI MIGQSHGSGAMFGALIASGIYVVLVSGIFSKVANLFPSIVTGSVITTIGLTLIPVAIGNMGNNVPEPTGQSLLLAAITVLIILLINIFTKG FIKSISILIGLVVGTAIAATMGLVDFSPVAVAPLVHVPTPLYFGMPTFEISSIVMMCIIATVSMVESTGVYLALSDITKDPIDSTRLRNG YRAEGLAVLLGGIFNTFPYTGFSQNVGLVKLSGIKKRLPIYYAAGFLVLLGLLPKFGALAQIIPSSVLGGAMLVMFGFVSIQGMQIL ARVDFANNEHNFLIAAVSIAAGVGLNNSNLFVSMPTAFQMFFSNGIVVASLLAIVLNAVLNHKKKZ

(SEQ. ID. NO. 36)

MKDRIKEYLQDKGKVTVNDLAQALGKDSSKDFRELIKTLSLMERKHQIRFEEDGSLTLEIKKKHEITLKGIFHAHKNGFGFVSLEGE EDDLFVGKNDVNYAIDGDTVEVVIKKVADRNKGTAAEAKIIDILEHSLTTVVGQIVLDQEKPKYAGYIRSKNQKISQPIYVKKPAL KLEGTEVLKVFIDKYPSKKHDFFVASVLDVVGHSTDVGIDVLEVLESMDIVSEFPEAVVKEAESVPDAPSQKDMEGRLDLRDEITF TIDGADAKDLDDAVHIKALKNGNLEFGVHIADVSYYVTEGSALDKEALNRATSVYVTDRVVPMLPERLSNGICSLNPQVDRLTQS AIMEIDKHGRVVNYTITQTVIKTSFRMTYSDVNDILAGDEEKRKEYHKIVSSIELMAKLHETLENMRVKRGALNFDTNEAKILVDK QGKPVDIVLRQRGIAERMIESFMLMANETVAEHFSKLDLPFIYRIHEEPKAEKVQKFIDYASSFGLRIYGTASEISQEALQDIMRAVE GEPYADVLSMMLLRSMQQARYSEHNHGHYGLAADYYTHFTSPIRRYPDLLVHRMIRDYGRSKEIAEHFEQVIPEIATQSSNRERRA EAEREVEAMKKAEYMEEYVGEEYDAVVSSIVKFGLFVELPNTVEGLIHITNLPEFYHFNERDLTLRGEKSGITFRVGQQIRIRVERA DKMTGEIDFSFVPSEFDVIEKGLKQSSRSGRGRDSNRRSDKKEDKRKKSGRSNDKRKHSQKDKKKKGKKFFYKEVAKKGAKHGKG RGKGRTKZ

(SEQ. ID. NO. 37)

MGTTGFTIIDLIILIVYLLAVLVAGIYFSKKEMKGKEFFKGDGSVPWYVTSVSIFATMLSPISFLGLAGSSYAGSWILWFAQLGMVV AIPLTIRFILPIFARIDIDTAYDYLDKRFNSKALRIISALLFIIYQLGRMSIIMYLPSAGLSVLTGIDINILIILMGVVAIVYSYTGGLKSVL WTDFIQGVILISGVVLALFVLIANIKGGFGAVAETLANGKFLAANEKLFDPNLLSNSIFLIVMGSGFTILSSYASSQDLVQRFTTTQNI KKLNKMLFTNGVLSLATATVFYLIGTGLYVFYQVQNADSAASNIPQDQIFMYFIAYQLPVGITGLILAAIYAASQSTISTGLNSVATS WTLDIQDVISKNMSDNRRTKIAQFVSLAVGLFSIGVSIVMAHSDIKSAYEWFNSFMGLVLGLLGGVFILGFVSKKANKQGAYAALI VSTIVMVFIKYFLPPTAVSYWAYSLISISVSVVSGYIVSVLTGNKVSAPKYTTIHDITEIKADSSWEVRHZ

(SEQ. ID. NO. 38)

MKFSKKYIAAGSAVIVSLSLCAYALNQHRSQENKDNNRVSYVDGSQSSQKSENLTPDQVSQKEGIQAEQIVIKITDQGYVTSHGDH YHYYNGKVPYDALFSEELLMKDPNYQLKDADIVNEVKGGYIIKVDGKYYVYLKDAAHADNVRTKDEINRQKQEHVKDNEKVNS NVAVARSQGRYTTNDGYVFNPADIIEDTGNAYIVPHGGHYHYIPKSDLSASELAAAKAHLAGKNMQPSQLSYSSTASDNNTQSVA KGSTSKPANKSENLQSLLKELYDSPSAQRYSESDGLVFDPAKIISRTPNGVAIPHGDHYHFIPYSKLSALEEKIARMVPISGTGSTVST NAKPNEVVSSLGSLSSNPSSLTTSKELSSASDGYIFNPKDIVEETATAYIVRHGDHFHYIPKSNQIGQPTLPNNSLATPSPSLPINPGTS HEKHEEDGYGFDANRIIAEDESGFVMSHGDHNHYFFKKDLTEEOIKVRKNIZ

(SEQ.JD, NO. 39)

MKKRAIVAVIVLLLIGLDQLVKSYIVQQIPLGEVRSWIPNFVSLTYLQNRGAAFSILQDQQLLFAVITLVVVIGAIWYLHKHMEDSF WMVLGLTLIIAGGLGNFIDRVSQGFVVDMFHLDFINFAIFNVADSYLTVGVIILLIAMLKEEINGNZ

(SEQ. ID. NO. 40)

MNTNLASFIVGLIIDENDRFYFVQKDGQTYALAKEEGQHTVGDTVKGFAYTDMKQKLRLTTLEVTATQDQFGWGRVTEVRKDLG VFVDTGLPDKEIVVSLDILPELKELWPKKGDQLYIRLEVDKKDRIWGLLAYQEDFQRLARPAYNNMQNQNWPAIVYRLKLSGTFV YLPENNMLGFIHPSERYAEPRLGQVLDARVIGFREVDRTLNLSLKPRSFEMLENDAQMILTYLESNGGFMTLNDKSSPDDIKATFGI SKGQFKKALGGLMKAGKIKQDQFGTELIZ

(SEQ. ID. NO. 41)

MKDVSLFLLKKVFKSRLNWIVLALFVSVLGVTFYLNSQTANSHSLESRIAANERAINENEEKLSQMSDTSSEEYQFAKNNLD VQKNLLTRKTEILTLLKEGRWKEAYYLQWQDEEKNYEFVSNDPTASPGLKMGVDRERKIYQALYPLNIKAHTLEFPTHGIDQIVWI LEVIIPSLFVVAIIFMLTQLFAERYQNHLDTAHLYPVSKVTFAISSLGVGVGYVTVLFIGICGFSFLVGSLISGFGQLDYPYPIYSLVNQ EVTIGKIQDVLFPGLLAFLAFIVIVEVVYLIAYFFKQKMPVLFLSLIGIVGLLFGIQTIQPLQRIAHLIPFTYLRSVEILSGRLPKQIDNV DLNWSMGMVLLPCLIIFLLLGILFIERWGSSQKKEFFNRFZ

(SEQ. ID. NO. 42)

MMKFILDIVSTPAILVALIAILGLVLQKKKLPDIIKGGIKTFVGFLVVSGGAGIVQNSLNPFGTMFEHAFHLSGVVPNNEAIVAVALT TYGSATAMIMFAGMVFNILIARFTRFKYIFLTGHHTLYMACMIAVILSVAGFTSLPLILLGGLALGIIMSISPAFVQKYMVQLTGNDK VALGHFSSLGYWLSGFTGSLIGDKSKSTEDIKFPKSLAFLRDSTVSITLSMAVIYIIVAIFAGSEYIEKEISSGTSGLVYALQLAGQFAA GVFVILAGVRLILGEIVPAFKGISERLVPNSKPALDCPIVYTYAPNAVLIGFISSFVGGLVSMVIMIASGTVVILPGVVPHFFCGATAG VIGNASGGVRGATIGAFLQGILISFLPVFLMPVLGGLGFQGSTFSDADFGLSGIILGMLNQFGSQAGIVIGLVLILAVMFGVSFIKKPS ATEEZ

(SEQ. ID. NO. 43)

MIKTFLSALSVILFSIPIITYSFFPSSNLNIWLSTQPILAQIYAFPLATATMAAILSFLFFFLSFYKKNKQIRFYSGILLLLSLILLLFGTDK TLSSASNKTKTLKLVTWNVANQIEAQHIERIFSHFDADMAIFPELATNIRGEQENQRIKLLFHQVGLSMANYDIFTSPPTNSGIAPVT VIVKKSYGFYTEAKTFHTTRFGTIVLHSRKQNIPDIIALHTAPPLPGLMEIWKQDLNIIHNQLASKYPKAIIAGDFNATMRHGALAKI SSHRDALNALPPFERGTWNSQSPKLFNATIDHILLPKNHYYVKDLDIVSFQNSDHRCIFTEITFZ

(SEQ. ID. NO. 44)

MNPIQRSWAYVSRKRLRSFILFLILLVLLAGISACLTLMKSNKTVESNLYKSLNTSFSIKKIENGQTFKLSDLASVSKIKGLENVSPEL ETVAKLKDKEAVTGEQSVERDDLSAADNNLVSLTALEDSSKDVTFTSSAFNLKEGRHLQKGDSKKILIHEELAKKNGLSLHDKIGL DAGQSESGKGQTVEFEIIGIFSGKKQEKFTGLSSDFSENQVFTDYESSQTLLGNSEAQVSAARFYVENPKEMDGLMKQVENLALEN QGYQVEKENKAFEQIKDSVATFQTFLTIFLYGMLIAGAGALILVLSLWLRERVYEVGILLALGKGKSSIFLQFCLEVVLVSLGALLP AFVAGNAITTYLLQTLLASGDQASLQDTLAKASSLSTSILSFAESYVFLVLLSCLSVALCFLFRKSPKEILSSISZ

(SEQ. ID. NO. 45)

MLHNAFA YVTRKFFKSIVIFLIILLMASLSLVGLSIKGATAKASQETFKNITNSFSMQINRRVNQGTPRGAGNIKGEDIKKITENKAIE SYVKRINAIGDLTGYDLIETPETKKNLTADRAKRFGSSLMITGVNDSSKEDKFVSGSYKLVEGEHLTNDDKDKILLHKDLAAKHG WKVGDKVKLDSNIYDADNEKGAKETVEVTIKGLFDGHNKSAVTYSQELYENTAITDIHTAAKLYGYTEDTAIYGDATFFVTADK NLDDVMKELNGISGINWKSYTLVKSSSNYPALEQSISGMYKMANLLFWGSLSFSVLLLALLLSLWINARRKEVGILLSIGLKQASIL GQFITESILIAIPALVSAYFLANYTARAIGNTVLANVTSGVAKQASKAAQASNLGGGAEVDGFSKTLSSLDISIQTSDFIIIFVLALVLV VLVMALASSNLLRKQPKELLLDGEZ

(SEQ. ID. NO. 46)

MSQDKQMKAVSPLLQRVINISSIVGGVGSLIFCIWAYQAGILQSKETLSAFIQQAGIWGPPLFIFLQILQTVVPIIPGALTSVAGVFIYGHIIGTIYNYIGIVIGCAIIFYLVRLYGAAFVQSVVSKRTYDKYIDWLDKGNRFDRFFIFMMIWPISPADFLCMLAALTKMSFKRYMTIIILTKPFTLVVYTYGLTYIIDFFWQMLZ

(SEQ. ID. NO. 47)

MRNMWVVIKETYLRHVESWSFFFMVISPFLFLGISVGIGHLQGSSMAKNNKVAVVTTVPSVAEGLKNVNGVNFDYKDEASAKEAI KEEKLKGYLTIDQEDSVLKAVYHGETSLENGIKFEVTGTLNELQNQLNRSTASLSQEQEKRLAQTIQFTEKIDEAKENKKFIQTIAA GALGFFLYMILITYAGVTAQEVASEKGTKIMEVVFSSIRASHYFYARMMALFLVILTHIGIYVVGGLAAVLLFKDLPFLAQSGILDH LGDAISLNTLLFILISLFMYVVLAAFLGSMVSRPEDSGKALSPLMILIMGGFFGVTALGAAGDNLLLKIGSYIPFISTFFMPFRTINDY AGGAEAWISLAITVIFAVVATGFIGRMYASLVLQTDDLGIWKTFKRALSYKZ

(SEQ. ID. NO. 48)

MTETIKLMKAHTSVRRFKEQEIPQVDLNEILTAAQMASSWKNFQSYSVIVVRSQEKKDALYELVPQEAIRQSAVFLLFVGDLNRAE KGARLHTDTFQPQGVEGLLISSVDAALAGQNALLAAESLGYGGVIIGLVRYKSEEVAELFNLPDYTYSVFGMALGVPNQHHDMKP

RLPLENVVFEEEYQEQSTEAIQAYDRVQADYAGARATTSWSQRLAEQFGQAEPSSTRKNLEQKKLLZMLKLIAIVGTNSKRSTNR QLLQYMQKHFTDKAEIELVEIKAIPVFNKPADKQVPAEILEIAAKIEEADGVIIGTPEYDHSIPAVLMSALAWLSYGIYPLLNKPIMIT GASYGTLGSSRAQLQLRQILNAPEIKANVLPDEFLLSHSLQAFNPSGDLVDLDVIKKLDAIFDDFRIFVKITEKLRNAQELLRKDAED FDWENLZ

(SEQ. ID. NO. 49)

MNTYQLNNGVEIPVLGFGTFKAKDGEEAYRAVLEALKAGYRHIDTAAIYQNEESVGQAIKDSGVPREEMFVTTKLWNSQQTYEQ TRQALEKSIEKLGLDYLDLYLIHWPNPKPLRENDAWKTRNAEVWRAMEDLYQEGKIRAIGVSNFLPHHLDALLETATIVPAVNQV RLAPGVYQDQVVAYCREKGILLEAWGPFGQGELFDSKQVQEIAANHGKSVAQIALAWSLAEGFLPLPKSVTTSRIQANLDCFGIEL SHEERETLKTIAVQSGAPRVDDVDFZ

(SEO, ID, NO, 50)

MRCKMLDPIAIQLGPLAIRWYALCIVTGLILAVYLTMKEAPRKKIIPDDILDFILVAFPLAILGARLYYVIFRFDYYSQNLGEIFAIWN GGLAIYGGLITGALVLYIFADRKLINTWDFLDIAAPSVMIAQSLGRWGNFFNQEAYGATVDNLDYLPGFIRDQMYIEGSYRQPTFL YESLWNLLGFALILIFRRKWKSLRRGHITAFYLIWYGFGRMVIEGMRTDSLMFFGFRVSQWLSVVLIGLGIMIVIYQNRKKAPYYIT EEENZ

(SEQ. ID. NO. 51)

MGKLSSILLGTVSGAALALFLTSDKGKQVCSQAQDFLDDLREDPEYAKEQVCEKLTEVKEQATDFVLKTKEQVESGEITVDSILAQ TKSYAFQATEASKNQLNNLKEQWQEKAEALDDSEEIVIDITEEZ

(SEQ. ID. NO. 52)

MKTKLIFWGSMLFLLSLSILLTIYLAWIFYPMEIQWLNLTNRVYLKPETIQYNFHILMNYLTNPFSQVLQMPDFRSSAAGLHHFAVV KNLFHLVQLVALVTLPSFYVFVNRIVKKDFLSLYRKSLLALVVLPVMIGLGGVLIGFDQFFTLFHQILFVGDDTWLFDPAKDPVIMI LPETFFLHAFLLFFALYENFFGYLYLKSRRKZ

(SEQ. ID. NO. 53)

MTYHFTEEYDIIVIGAGHAGVEASLAASRMGCKVLLATINIEMLAFMPCNPSIGGSAKGIVVREVDALGGEMAKTIDKTYIQMKML NTGKGPAVRALRAQADKELYSKEMRKTVENQENLTLRQTMIDEILVEDGKVVGVRTATHQEYAAKAVIVTTGTALRGEIIIGDLK YSSGPNHSLASINLADNLKELGLEIGRFKTGTPPRVKASSINYDVTEIQPGDEVPNHFSYTSRDEDYVKDQVPCWLTYTNGTSHEIIQ NNLHRAPMFTGVVKGVGPRYCPSIEDKIVRFADKERHQLFLEPEGRNTEEVYVQGLSTSLPEDVQRDLVHSIKGLENAEMMRTGY AIEYDMVLPHQLRATLETKKISGLFTAGQTNGTSGYEEAAGQGIIAGINAALKIQGKPELILKRSDGYIGVMIDDLVTKGTIEPYRLL TSRAEYRLILRHDNADMRLTEMGREIGLVDDERWARFEIKKNQFDNEMKRLDSIKLKPVKETNAKVEEMGFKPLTDAVTAKEFLR RPEVSYQDVVAFIGPAAEDLDDKIIELIETEIKYEGYISKAMDQVAKMKRMEEKRIPANIDWDDIDSIATEARQKFKLINPETIGQAS RISGVNPADISILMVYLEGKNRSISKTLOKSKZ

(SEQ. ID. NO. 54)

MTKQVLLVDDEEHILKLLDYHLSKEGFSTQLVTNGRKALALAETEPFDFILLDIMLPQLDGMEVCKRLRAKGVKTPIMMVSAKSD EFDKVLALELGADDYLTKPFSPRELLARVKAVLRRTKGEQEGDDSDNIADDSWLFGTLKVYPERHEVYKANKLLSLTPKEFESDK NPFFEVFKVSKVTAQZ

(SEQ. ID. NO. 55)

MTTFKDGFLWGGAVAAHQLEGGWQEGGKGISVADVMTAGRHGVAREITLGVLEGKYYPNHEAIDFYHRYKEDIALFAEMGFKC FRTSIAWTRIFPKGDELEPNEEGLQFYDNLFDECLKNGIEPVITLSHFEMPYHLVTEYGGWKNRKLIDFFARFAEVVFKRYKDKVK YWMTFNEINNQANYQEDFAPFTNSGIVYEEGDNREAIMYQAAHYELVASARAVKIGHEINPDFQIGCMIAMCPIYPVTCNPKDILM AMKAMQKRYYFADVHVLGKYPEHIFKYWERKGISVDFTAQDKEDLLGGTVDYIGFSYYMSFAIDSHRENNPYFDYLETEDLVKN NYVKASEWEWQIDPEGLRYALNWFTDHYHLPLFIVENGFGAIDQVAADGMVHDDYRIEYLGAHIREMKKAVVEDGVDLMGYTP WGCIDLVSAGTGEMRKRYGFIYVDKDDNGKGSYNRSPKKSFGWYKEVISSNGESVEZ

(SEQ. ID. NO. 56)

MDQQNGLFGFLENHVMGPMGKLAQFKVVRAITAAGMAAVPFTIVGSMFLVFSILPQAFSFWPIVADIFSASFDKFTSLYMVANYA TMGSLSLYFVLSLAYELTKIYAEEEELNMNPLNGALLALMAFVMTVPQIIFDGGMMKTVTSLKEGAVIADGWAMGNVVARFGTT GIFTAIIMAIVTVLIYRMCVKHNWVIKMPEAVPEGVSRGFTALVPGFVVAFVVIFINGLLVAMGTDIFKVIAIPFGFVSNLTNSWIGL MIIYLLTQLLWIVGIHGANIVFAFVSPIALANMAENAAGGHFAVAGEFSNMFVIAGGSGATLGLCLYIAFASKSEQLKAIGRASVVP ALFNINEPLIFGLPIIYNPALAIPFILAPMVTATIYYVANSLNFIKPIIAQVPWPTPVGIGAFLGTADLRAVLVALVCAFAAFLVYLPFIR VYDQKLVKEEQGIZ

(SEO. ID. NO. 57)

MKKFYVSPIFPILVGLIAFGVLSTFIIFVNNNLLTVLILFLFVGGYVFLFKKLRVHYTRSDVEQIQYVNHQAEESLTALLEQMPVGVM KLNLSSGEVEWFNPYAELILTKEDGDFDLEAVQTIIKASVGNPSTYAKLGEKRYAVHMDASSGVLYFVDVSREQAITDELVTSRPV IGIVSVDNYDDLEDETSESDISQINSFVANFISEFSEKHMMFSRRVSMDRFYLFTDYTVLEGLMNDKFSVIDAFREESKQRQLPLTLS MGFSYGDGNHDEIGKVALLNLNAEVRGGDQVVVKENDETKNPVYFGGGSAASIKRTRTRAMMTAISDKIRSVDQVFVVGHK NLDMDALGSAVGMQLFASNVIENSYALYDEEQMSPDERAVSFIEKEGVTKLLSVKDAMGMVTNRSLLILVDHSKTALTLSKEFY DLFTQTIVIDHHRRDQDFPDNAVITYIESGASSASELVTELIQFQNSKKNRLSRMQASVLMAGMMLDTKNFTSRVTSRTFDVASYL RTRGSDSIAIQEIAATDFEEYREVNELILQGRKLGSDVLIAEAKDMKCYDTVVISKAADAMLAMSGIEASFVLAKNTQGFISISARSR SKLNVQRIMEELGGGGHFNLAAAQIKDVTLSEAGEKLTEIVLNEMKEKEKEEZ

(SEQ. ID. NO. 58)

MKEKNMWKELLNRAGWILVFLLAVLLYQVPLVVTSILTLKEVALLQSGLIVAGLSIVVLALFIMGARKTKLASFNFSFFRAKDLAR LGLSYLVIVGSNILGSILLQLSNETTTANQSQINDMVQNSSLISSFFLLALLAPICEEILCRGIVPKKIFRGKENLGFVVGTIVFALLHQP SNLPSLLIYGGMSTVLSWTAYKTQRLEMSILLHMIVNGIAFCLLALVVIMSRTLGISVZ

(SEQ. ID. NO. 59)

 ${\tt MKEKNMWKELLNRAGWILVFLLAVLLYQVPLVVTSILTLKEVALLQSGLIVAGLSIVVLALFIMGARKTKLASFNFSFRAKDLAR}$ LGLSYLVIVGSNILGSILLQLSNETTTANQSQINDMVQNSSLISSFFLLALLAPICEEILCRGIVPKKIFRGKENLGFVVGTIVFALLHQP SNLPSLLIYGGMSTVLSWTAYKTORLEMSILLHMIVNGIAFCLLALVVIMSRTLGISVZ

(SEQ. ID. NO. 60)

MDTQKIEAAVKMIIEAVGEDANREGLQETPARVARMYQEIFSGLGQTAEEHLSKSFEIIDDNMVVEKDIFFHTMCEHHFLPFYGRA HIAYIPDGRVAGLSKLARTVEVYSKKPQIQERLNIEVADALMDYLGAKGAFVVIEAEHMCMSMRGVRKPGTATLTTVARGLFETD KDLRDQAYRLMGLZMKDLFLKRKQAFRKECLGYLRYVLNDHFVLFLLVLLGFLAYQYSQLLQHFPENHWPILLFVGITSVLLLLW GGTATYMEAPDKLFLLVGEEEIKLHLKRQTGISLVFWLFVQTLFLLLFAPLFLAMGYGLPVFLLYVLLLGVGKYFHFCQKASKFFT ETGLDWDYVISQESKRKQVLLRFFALFTQVKGISNSVKRRAYLDFILKAVQKVPGKIWQNLYLRSYLRNGDLFALSLRLLLLSLLA OVFIEQAWIATAVVVLFNYLLLFQLLALYHAFDYQYLTQLFPLDKGQKEKGLQEVVRGLTSFVLLVELVVGLITFQEKLALLALLG AGLVLLVLYLPYQVKRQMQDZ

(SEQ. ID. NO. 61)

MRKSIVLAADNAYLIPLETTIKSVLYHNRDVDFYILNSDIAPEWFKLLGRKMEVVNSTIRSVHIDKELFESYKTGPHINYASYFRFFA TEVVESDRVLYLDSDIIVTGELATLFEIDLKGYSIGAVDDVYAYEGRKSGFNTGMLLMDVAKWKEHSIVNSLLELAAEONOVVHL GDQSILNIYFEDNWLALDKTYNYMVGIDIYHLAQECERLDDNPPTIVHYASHDKPWNTYSISRLRELWWYYRDLDWSEIAFORSD LNYFERSNQSKKQVMLVTWSADIKHLEYLVQRLPDWHFHLAAPCDCSEELTSLSQYTNVTVYQNVLHSRIDWLLDDSIVYLDINT GGEVFNVVTRAQESGKKIFAFDITRKSMDDGLYDGIFSVERPDDLVDRMKNIEIEZ

(SEQ. ID. NO. 62)

MTKIYSSIAVKKGLFTSFLLFIYVLGSRIILPFVDLNTKDFLGGSTAYLAFSAALTGGNLRSLSIFSVGLSPWMSAMILWQMFSFSKRL GLTSTSIEIQDRRKMYLTLLIAVIQSLAVSLRLPVQSSYSAILVVLMNTILLIAGTFFLVWLSDLNASMGIGGSIVILLSSMVLNIPQDV LETFQTVHIPTGIIVLLALLTLVFSYLLALMYRARYLVPVNKIGLHNRFKRYSYLEIMLNPAGGMPYMYVMSFLSVPAYLFILLGFIF PNHSGLAALSKEFMVGKPLWVYVYISVLFLFSIIFAFVTMNGEEIADRMKKSGEYIYGIYPGADTSRFINRLVLRFSVIGGLFNVIMA GGPMLFVLFDEKLLRLAMIPGLFMMFGGMIFTIRDEVKALRLNETYRPLIZ

(SEQ. ID. NO. 63)

MSSLSDQELVAKTVEFRQRLSEGESLDDILVEAFAVVREADKRILGMFPYDVQVMGAIVMHYGNVAEMNTGEGKTLTATMPVYL NAFSGEGVMVVTPNEYLSKRDAEEMGQVYRFLGLTIGVPFTEDPKKEMKAEEKKLIYASDIIYTTNSNLGFDYLNDNLASNEEGKF LRPFNYVIIDEIDDILLDSAQTPLIIAGSPRVQSNYYAIIDTLVTTLVEGEDYIFKEEKEEVWLTTKGAKSAENFLGIDNLYKEEHASF ARHLVYAIRAHKLFTKDKDYIIRGNEMVLVDKGTGRLMEMTKLQGGLHQAIEAKEHVKLSPETRAMASITYQSLFKMFNKISGMT GTGKVAEKEFIETYNMSVVRIPTNRPRQRIDYPDNLYITLPEKVYASLEYIKQYHAKGNPLLVFVGSVEMSQLYSSLLFREGIAHNV LNANNAAREAQIISESGQMGAVTVATSMAGRGTDIKLGKGVAELGGLIVIGTERMESQRIDLQIRGRSGRQGDPGMSKFFVSLEDD VIKKFGPSWVHKKYKDYQVQDMTQPEVLKGRKYRKLVEKAQHASDSAGRSARRQTLEYAESMNIQRDIVYKERNRLIDGSRDLE DVVVDIIERYTEEVAADHYASRELLFHFIVTNISFHVKEVPDYIDVTDKTAVRSFMKQVIDKELSEKKELLNQHDLYEQFLRLSLLK AIDDNWVEQVDYLQQLSMAIGGQSASQKNPIVEYYQEAYAGFEAMKEQIHADMVRNLLMGLVEVTPKGEIVTHFPZ

(SEQ. ID. NO. 64)

MIGTFAAALVAVLANFIVPIEITPNSANTEIAPPDGIGQVLSNLLLKLVDNPVNALLTANYIRILSWAVIFGIAMREASKNSQELLKTI ADVTSKIVEWIINLAPFGILGLVFKTISDKGVGSLANYGILLVLLVTTMLFVAPVVNPLIAFFFMRRNPYPLVWNCLRVSGVTAFFTR SSATNIPVNMKLCHDLGLNPDTYSVSIPLGSTINMAGVAITINLLTLAAVNTLGIPVDFATAFVLSVVAAISSCDASGIAGGSLLLIPV ACSLFGISNDIAIQIVGVGFVIGVIQDSCETALNSSTDVLFTAVAEYAATRKKZ

MSISQRTTKLILATCLACLLAYFLNLSSAVSAGIIALLSLSDTRRSTLKLARNRLFSMLLALAIGVLAFHLSGFHIWSLGLYLAFYVPL AYKMGWEIGITPSTVLVSHLLVQESTSPDLLVNEFLLFAIGTGFALLVNLYMPSREEEIQHYHTLVEEKLKDILQRFKYYLSRGDGR NRAQLVAELDTLLKEALRLVYLDHSDHLFHQTDYHIHYFEMRQRQSRILRNMAQQINTCHLAASESLILAQLFSKIAGQLSQTNPA SDLLDEIERYLEVFRNRSLPKTREEFETRATLLQLLREAKTFIQVKVDFYQKYRQZ

(SEQ. ID. NO. 66)
MEIMSLAIAVFAVIIGLVIGYVSISAKMKSSQEAAELMLLNAEQEATNLRGQAEREADLLVNEAKRESKSLKKEALLEAKEEARKY REEVDAEFKSERQELKQIESRLTERATSLDRKDDNLTSKEQTLEQKEQSISDRAKNLDAREEQLEEVEROKEAELERIGALSOAEAR DIILAOTEENLTREIASRIREAEQEVKERSDKMAKDILVQAMQRIAGEYVAESTNSTVHLPDDTMKGRIIGREGRNIRTFESLTGVDV IIDDTPEVVTLSGFDPIRREIARMTMEMLLKDGRIHPARIEELVEKNRQEIDNKIREYGEAAAYEIGAPNLHPDLMKIMGRLQFRTSY GQNVLRHSIEVAKLAGIMASELGENAALARRAGFLHDIGKAIDHEVEGSHVEIGMELARKYKEPPVVVNTIASHHGDVEAESVIAV IVAAADALSAARPGARSESLESYIKRLHDLEEIANGFEGVQTSFALQAGREIRIMVNPGKIKDDKVTILAHKVRKKIENNLDYPGNI KVTVIRELRAVDYAKZ

MMLKPSIDTLLDKVPSKYSLVILEAKRAHELEAGAPATQGFKSEKSTLRALEEIESGNVTIHPDPEGKREAVRRRIEEEKRRKEEEEK KIKEQIAKEKEDGEKIZ

(SEQ. ID. NO. 68)

MSAYOLPTVWQDEASNQGAFTGLNRPTAGARFEQNLPKGEQAFQLYSLGTPNGVKVTILLEELLEAGFKEAAYDLYKIAIMDGDQ FGSDFVKLNPNSKIPALLDQSGTENVRVFESAHILLYLAEKFGAFLPSNPVEKVEVLNWLFWQAGAAPFLGGGFGHFFNYAPEKLE YPINRFTMEVKRQLDLLDKELAQKPYIAGNDYTIADIAIWSWYGQLVQGNLYQGSAKFLDASSYQNLVKWAEKIANRPAVKRGL **EVTYTEIKZ**

(SEQ. ID. NO. 69)

LASLITSIIMFYVGFDVLRDTIQKILSREETVIDPLGATLGIISAAIMFVVYLYNTRLSKKSNSNALKAAAKDNLSDAVTSLGTAIAILA SSFNYPIVDKLVAIIITFFILKTAYDIFIESSFSLSDGFDDRLLEDYQKAIMEIPKISKVKSQRGRTYGSNIYLDITLEMNPDLSVFESHEI ADQVESMLEERFGVFDTDVHIEPAPIPEDEILDNVYKKLLMREQLIDQGNQLEELLTDDFVYIRQDGEQMDKEAYKTKKELNSAIK DIQITSISQKTKLICYELDGIIHTSIWRRHETWONIFHOETKKEZ

(SEQ. ID. NO. 70)

MTIKLVATDMDGTFLDGNGRFDMDRLKSLLVSYKEKGIYFAVASGRGFLSLEKLFAGVRDDIIFIAENGSLVEYQGQDLYEATMS RDFYLATFEKLKTSPYVDINKLLLTGKKGSYVLDTVDETYLKVSQHYNENIQKVASLEDITDDIFKFTTNFTEETLEDGEAWVNEN VPGVKAMTTGFESIDIVLDYVDKGVAIVELVKKLGITMDQVMAFGDNLNDLHMMQVVGHPVAPENARPEILELAKTVIGHHKER SVIAYMEGLZ

(SEQ. ID. NO. 71)

MADIKLIALDLDGTLLTTDKRLTDRTKETLQAARDRGIKVVLTTGRPLKAMDFFLHELGTDGQEDEYTITFNGGLVQKNTGEILDK TVFSYDDVARLYEETEKLSLPLDAISEGTVYOIOSDOESLYAKFNPALTFVPVDFEDLSSOMTYNKCVTAFAOEPLDAAIOKISPELF DQYEIFKSREMLLEWSPKNVHKATGLAKLISHLGIDQSQVMACGDEANDLSMIEWAGLGVAMQNAVPEVKAAANVVTPMTNDE EAVAWAIEEYVLKENZ

(<u>SEQ. ID. NO. 72)</u> MESLLILLIANLAGLFLIWQRQDRQEKHLSKSLEDQADHLSDQLDYRFDQARQASQLDQKDLEVVVSDRLQEVRIELHQGLTQV RQEMTDNLLQTRDKTDQRLQALQESNEQRLEQMRQTVEEKLEKTLQTRLQASFETVSKQLESVNRGLGEMQTVARDVGALNKV LSGTKTRGILGELOLGOIJEDIMTPAOYEREYATVENSSERVEYAJKLPGOGDOEYVYLPJDSKFPLADYYRLEFAYETGDKDEJERC RKSLLASVKRFARDIRNKYIAPPRTTNFGVLFVPTEGLYSEIVRNPVFFDDLRREEQIIVAGPSTLSALLNSLSVGFKTLNIQKSADHIS KTLASVKTEFGKFGGILVKAQKHLQHASGNIDELLNRRTIAIERTLRHIELSEGEPALDLLHFOENEEEYEDZ

(<u>SEQ. ID. NO. 73)</u> MKISHMKKDELFEGFYLIKSADLRQTRAGKNYLAFTFQDDSGEIDGKLWDAQPHNIEAFTAGKVVHMKGRREVYNNTPQVNQIT LRLPQAGEPNDPADFKVKSPVDVKEIRDYMSQMIFKIENPVWQRIVRNLYTKYDKEFYSYPAAKTNHHAFETGLAYHTATMVRL ADAISEVYPOLNKSLLYAGIMLHDLAKVIELTGPDOTEYTVRGNLLGHIALIDSEITKTVMELGIDDTKEEVVLLRHVII.SHHGLLE YGSPVRPRIMEAEIIHMIDNLDASMMMMSTALALVDKGEMTNKIFAMDNRSFYKPDLDZ

(SEO. ID. NO. 74)

MSEKAKKGFKMPSSYTVLLIIIAIMAVLTWFIPAGAFIEGIYETQPQNPQGIWDVLMAPIRAMLGTHPEEGSLIKETSAAIDVAFFILM VGGFLGIVNKTGALDVGIASIVKKYKGREKMLILVLMPLFALGGTTYGMGEETMAFYPLLVPVMMAVGFDSLTGVAIILLGSQIGC LASTLNPFATGIASATAGVGTGDGIVLRLIFWVTLTALSTWFVYRYADKIQKDPTKSLVYSTRKEDLKHFNVEESSSVESTLSSKQK SVLFLFVLTFILMVLSFIPWTDLGVTIFDDFNTWLTGLPVIGNIVGSSTSALGTWYFPEGAMLFAFMGILIGVIYGLKEDKIISSFMNG AADLLSVALIVAIARGIQVIMNDGMITDTILNWGKEGLSGLSSQVFIVVTYIFYLPMSFLIPSSSGLASATMGIMAPLGEFVNVRPSLII TAYOSASGVLNLIAPTSGIVMGALALGRINIGTWWKFMGKLVVAIIVVTIALLLLGTFLPFLZ

MSNSFVKLLVSQLFANLADIFFRVTIIANIYIISKSVIATSLVPILIGISSFVASLLVPLVTKRLALNRVLSLSQFGKTILLAILVGMFTV MQSVAPLVTYLFVVAISILDGFAAPVSYAIVPRYATDLGKANSALSMTGEAVQLIGWGLGGLLFATIGLLPTTCINLVLYIISSFLML FLPNAEVEVLESETNLEILLKGWKLVARNPRLRLFVSANLLEIFSNTIWVSSIILVFVTELLNKTESYWGYSNTAYSIGIIISGLIAFRLS EKFLAAKWEPQLFTPNLKTIQNPCLSLDPGWFLFSPNGCFLLDKKEFPLYGISVEKNTKRKETHMNSLPNHHFONKSFYOLSFDGG HLTQYGGLIFFQELFSQLKLKERISKYLVTNDQRRYCRYSDSDILVQFLFQLLTGYGTDYACKELSADAYFPKLLEGGQLASQPTLS RFLSRTDEETVHSLRCLNLELVEFFLQFHQLNQLIVDIDSTHFTTYGKQEGVAYNAHYRAHGYHPLYAFEGKTGYCFNAQLRPGN RYCSEEADSFITPVLERFNQLLFRMDSGFATPKLYDLIEKTGQYYLIKLKKNTVLSRLGDLSLPCPODEDLTILPHSAYSETLYOAGS WSHKRRVCQFSERKEGNLFYDVISLVTNMTSGTSQDQFQLYRGRGQAENFIKEMKEGFFGDKTDSSTLIKNEVRMMMSCIAYNLY LFLKHLAGGDFQTLTIKRFRHLFLHVVGKCVRTGRKQLLKLSSLYAYSELFSALYSRIRKVNLNLPVPYEPPRRKASLMMHZ

MMEFFQQLPHLEPYGNPQYFVYVIAATLPIFIGLFFKKRFAWYEVLVSLFFIVTMLVGGKTNQLAALGIYLCWEILLLLFYKHYRKS KDGKWVFYLVSFLSLLPIIFVKVQPAINGTQSLLGFLGISYLTFRSVGIVIELRDGVIKDFTLWEFLRFLLFMPTFSSGPIDRFKRFNEN $YQAIPERDELMDMLDESVRYIM \hat{W}GFLYKFILAHVLGETLLPPLKNLALQSGGFFNLYALAVMYTFGLELFFDFAGYSMFALAISNL$ MGIRSPINFNKPFLSRDLKEFWNRWHMSLSFWFRDFVFMRMVMVLTRKKVFKNRNVTSSMAYIVNMLIMGFWHGVTWYYIAYG LFHGLGLVINDAWVRKKKTLNKERKKAGKAALPENRWIQLLGMVVTFHVVMLSFLIFSGFLNNLWFKKZ

(SEO. ID. NO. 77)

MLKRLWMIFGPVLIAGLLVFLLIFFYPTEMHHNLGAEKRSAVATTIDSFKERSQKVRALSDPNVRFVPFFGSSEWLRFDGAHPAVL AEKYNRSYRPYLLGOGGAASLNOYFGMQQMLPQLENKQVVYVISPQWFSKNGYDPAAFQQYFNGDQLTSFLKHQSGDQASQYA ATRLLQQFPNVAMKDLVQKLASKEELSTADNEMIELLARFNERQASFFGQFSVRGYVNYDKHVAKYLKILPDQFSYQAIEDVVKA DAEKNTSNNEMGMENYFYNEQIKKDLKKLKDSQKSFTYLKSPEYNDLQLVLTQFSKSKVNPIFIIPPVNKKWMNYAGLREDMYQ QTVQKIRYQLESQGFTNIADFSKDGGEPFFMKDTIHLGWLGWLAFDKAVDPFLSNPTPAPTYHLNERFFSKDWATYDGDVKEFQZ

(SEQ. ID. NO. 78)

MEKNLKALKQTTDQEGPAIEPEKAEDTKTVQNGYFEDAAVKDRTLSDYAGNWQSVYPFLEDGTFDQVFDYKAKLTGKMTQAEY KAYYTKGYHTDVTKINITDNTMEFVQGGQSKKYTYKYVGKKILTYKKGNRGVRFLFEATDADAGGFKYVQFSDHNVAPVKAEH FHIFFGGTSQEALFEEMDNWPTYYPDNLSGQEIAQEMLAHZ

(SEQ. ID. NO. 79)

MKDGHLLAHHIRLLNGRIFQKLLSODPEALYRGEQGKILAVLWNSETGCATATDIALATGLANNTLTTMIKKLEEOKLVIVSPCGK DKRKKYLVLTELGKSQKEVGHRVSQKLDTIFYKGFSEEEIHQFEGFQERILANLKEKGNEVZ

(SEQ. ID. NO. 80)

MTNLIATFQDRFSDWLTALSQHLQLSLLTLLLAILLAIPLAVFLRYHEKLADWVLQIAGIFQTIPSLALLGLFIPLMGIGTLPALTALV IYAIFPILQNTITGLKGIDPNLQEAGIAFGMTRWERLKKFEIPLAMPVIMSGIRTAAVLIIGTATLAALIGAGGLGSFILLGIDRNNASLI LIGALSSAVLAIAFNFLLKVMEKAKLRTIFSGFALVALLLGLSYSPALLVOKEKENLVIAGKIGPEPEILANMYKLLIEENTSMTATV KPNFGKTSFLYEALKKGDIDIYPEFTGTVTESLLQPSPKVSHEPEQVYQVARDGIAKQDHLAYLKPMSYQNTYAVAVPKKIAQEYG LKTISDLKKVEGQLKAGFTLEFNDREDGNKGLQSMYGLNLNVATIEPALRYQAIQSGDIQITDAYSTDAELERYDLQVLEDDKQLF PPYQGAPLMKEALLKKHPELERVLNTLAGKITESQMSQLNYQVGVEGKSAKQVAKEFLQEQGLLKKZ

(SEQ. ID, NO. 81)

MMHTYLQKKIENIKTTLGEMSGGYRRMVAAMADLGFSGTMKAIWDDLFAHRSFAQWIYLLVLGSFPLWLELVYEHRIVDWIGMI CSLTGIICVIFVSEGRASNYLFGLINSVIYLILALQKGFYGEVLTTLYFTVMQPIGLLVWIYQAQFKKEKQEFVARKLDGKGWTKYL SISVLWWLAFGFIYQSIGANRPYRDSITDATNGVGQILMTAVYREQWIFWAATNVFSIYLWWGESLQIQGKYLIYLINSLVGWYQ WSKAAKQNTDLLNZ

(SEQ. ID. NO. 82)

MRNMKAKYAVWVAFFLNLTYAIVEFIAGGVFGSSAVLADSVHDLGDAIAIGISAFLETISNREEDNQYTLGYKRFSLLGALVTAVI LVTGSVLVILENVTKILHPQPVNDEGILWLGIIAITINLLASLVVGKGKTKNESILSLHFLEDTLGWVAVILMAIVLRFTDWYILDPLL SLVISFFILSKALPRFWSTLKIFLDAVPEGLDIKQVKSGLERLDNVASLNQLNLWTMDALEKNAIVHVCLKEMEHMETCKESIRIFL KDCGFQNITIEIDADLETHOTHKRKVCDLERSYEHOHZ

(SEQ. ID. NO. 83)

MIEYKNVALRYTEKDVLRDVNLQIEDGEFMVLVGPSGSGKTTMLKMINRLLEPTDGNIYMDGKRIKDYDERELRLSTGYVLQAIA LFPNLTVAENIALIPEMKGWSKEEITKKTEELLAKVGLPVAEYGHRLPSELSGGEQQRVGIVRAMIGQPKIFLMDEPFSALDAISRK QLQVLTKELHKEFGMTTIFVTHDTDEALKLADRIAVLODGEIROVANPETILKAPATDFVADLFGGSVHDZ

(SEO, ID, NO, 84)

MSAVAISAMTKVMQETHGNPSSIHGHGRQAGKLLREARQELAQLLRTKPQHIFFTSGGTEGNNTTIIGYCLRHQEQGKHIITTAIEH HAVLETIDYLVQHFGFEATIIQPENQEITAQQIQKALRDDTILVSTMFVNNETGNLLPIAEIGQILKQHPAAYHVDAVQAIGKIPIHSE ELGIDFLTASAHKFHGPKGIGFLYASSMDFDSYLHGGDQEQKKRAGTENLPAIVGMVAALKEDLEKQEEHFQHVQNLETAFLAEL EGIQYYLNRGKHHLPYVLNIGFPGQKNDLLLLRLDLAGISISTGSACTAGVVQSSHVLEAMYGANSERLKESLRISLSPQNTVEDLQ TLAKTLKEIIGGZ

(SEQ. ID. NO. 85)

MLFKLSKEKIELGLSRLSPARRIFLSFALVILLGSLLLSLPFVQVESSRATYFDHLFTAVSAVCVTGLSTLPVAHTYNIWGQIICLLLIQ IGGLGLMTFIGVFYIQSKQKLSLRSRATIQDSFSYGETRSLRKFVYSIFLTTFLVESLGAILLSFRLIPQLGWGRGLFSSIFLAISAFCNA GFDNLGSTSLFAFQTDLLVNLVIAGLIITGGLGFMVWFDLAGHVGRKKKGRLHFHTKLVLLLTIGLLLFGTATTLFLEWNNAGTIG NLPVADKVLVSFFQTVTMRTAGFSTIDYTQAHPVTLLIYILQMFLGGAPGGTAGGLKITTFFVLLVFARSELLGLPHANVARRTIAP RTVQKSFSVFIIFLMSFLIGLILLGITAKGNPPFIHLVFETISALSTVGVTANLTPDLGKLALSVIMPLMFMGRIGPLTLFVSLADYHPE KKDMIHYMKADISIGZ

(SEQ. ID. NO. 86)

MSDRTIGILGLGIFGSSVLAALAKQDMNIIAIDDHAERINQFEPVLARGVIGDITDEELLRSAGIDTCDTVVVATGENLESSVLAVMH CKSLGVPTVIAKVKSQTAKKVLEKIGADSVISPEYEMGQSLAQTILFHNSVDVFQLDKNVSIVEMKIPQSWAGQSLSKLDLRGKYN LNILGFREQENSPLDVEFGPDDLLKADTYILAVINNQYLDTLVALNSZ

(SEQ. ID. NO. 87)

MKLLSIAISSYNAAAYLHYCVESLVIGGEQVGILIINDGSQDQTQEIAECLASKYPNIVRAIYQENKCHGGAVNRGLVEASGRYFKV VDSDDWVDPRAYLKILETLQELESKGQEVDVFVTNFVYEKEGQSRKKSMSYDSVLPVRQIFGWDQVGNFSKGQYTMMHSLIYRT DLLRASQFZ

(SEQ. ID. NO. 88)

MKFNPNQRYTR WSIRRLSVGVASVVVASGFFVLVGQPSSVRADGLNPTPGQVLPEETSGTKEGDLSEKPGDTVLTQAKPEGVTGN TNSLPTPTERTEVSEETSPSSLDTLFEKDEEAQKNPELTDVLKETVDTADVDGTQASPAETTPEQVKGGVKENTKDSIDVPAAYLEK AEGKGPFTAGVNQVIPYELFAGDGMLTRLLLKASDNAPWSDNGTAKNPALPPLEGLTKGKYFYEVDLNGNTVGKQGQALIDQLR ANGTQTYKATVKVYGNKDGKADLTNLVATKNVDININGLVAKETVQKAVADNVKDSIDVPAAYLEKAKGEGPFTAGVNHVIPY ELFAGDGMLTRLLLKASDKAPWSDNGDAKNPALSPLGENVKTKGQYFYQVALDGNVAGKEKQALIDQFRANGTQTYSATVNVYGNKDGKPDLDNIVATKKVTININGLISKETVQKAVADNVKDSIDVPAAYLEKAKGEGPFTAGVNHVIPYELFAGDGMLTRLLLKA SDKAPWSDNGDAKNPALSPLGENVKTKGQYFYQLALDGNVAGKEKQALIDQFRANGTQTYSATVNVYGNKDGKPDLDNIVATK KVTININGLISKETVQKAVADNVKTVSMFQQPTZ

(SEQ. ID. NO. 89)

MKLKSYILVGYIISTLLTILVVFWAVQKMLIAKGEIYFLLGMTIVASLVGAGISLFLLLPVFTSLGKLKEHAKRVAAKDFPSNLEVQG PVEFQQLGQTFNEMSHDLQVSFDSLEESEREKGLMIAQLSHDIKTPITSIQATVEGILDGIIKESEQAHYLATIGRQTERLNKLVEELN FLTLNTARNQVETTSKDSIFLDKLLIECMSEFQFLIEQERRDVHLQVIPESARIEGDYAKLSRILVNLVDNAFKYSAPGTKLEVVAKL EKDQLSISVTDEGQGIAPEDLENIFKRLYRVETSRNMKTGGHGLGLAIARELAHQLGGEITVSSQYGLGSTFTLVLNLSGSENKAZ

(SEQ. ID. NO. 90)

MFGQTAQHGLTNSLKDFWIFLLNIGPQLAFFCQMLRCSRSVEQGTGNHRREFNMIQQIFSHFGMTHLGQIKLVYQESIDLELLVNA LNHHLLIDRLVLTPNQITIEIDRQIVHGLDLLKGRKDKEIIDIKSMFRQLELASTQQICPINQRVHHGILAFGEISDLVPAKNLPNRQDZ

(SEQ. ID. NO. 91)

MEHLATYFSTYGGAFFAALGIVLAVGLSGMGSAYGVGKAGQSAAALLKEQPEKFASALILQLLPGTQGLYGFVIGILIWLQLTPEL PLEKGVAYFFVALPIAIVGYFSAKHQGNVAVAGMQILAKRPKEFMKGAILAAMVETYAILAFVVSFILTLRVZ

SEO. ID. NO. 92)

MLKSEKQSRYQMLNEELSFLLEGETNVLANLSNASALIKSRFPNTVFAGFYLFDGKELVLGPFQGGVSCIRIALGKGVCGEAAHFQ ETVIVGDVTTYLNYISCDSLAKSEIVVPMMKNGQLLGVLDLDSSEIEDYDAMDRDYLEQFVAILLEKTAWDFTMFEEKSZ

(SEQ. ID. NO. 93)

MSVLEIKDLHVEIEGKEILKGVNLTLKTGEIAAIMGPNGTGKSTLSAAIMGNPNYEVTKGEVLFDGVNILELEVDERARMGLFLAM QYPSEIPGITNAEFLRAAMNAGKEDDEKISVREFITKLDEKMELLNMKEEMAERYLNEGFSGGEKKRNEILQLLMLEPTFALLDEID SGLDIDALKVVSKGVNAMRGEGFGAMIITHYQRLLNYITPDVVHVMMEGRVVLSGGPELAARLEREGYAKLAEELGYDYKEELZ

(SEQ. ID. NO. 94)

MPYKRORSFSMALSKLDSLYMAVVADHSKNPHHQGKLEDAEQISLNNPTCGDVINLSVKFDAEDRLEDIAFLNSGCTISTASASM MTDAVLGKTKQEILELATIFSEMVQGQKDERQDQLGDAAFLSGVAKFPORIKCATLAWNALKKTIENOEKOZ

<u>(SEQ. ID. NO. 95)</u> MKIQDLLRKDVMLLDLQATEKTAVIDEMIKNLTDHGYVTDFETFKEGILAREALTSTGLGDGIAMPHSKNAAVKEATVLFAKSNK GVDYESLDGQATDLFFMIAAPEGANDTHLAALAELSQYLMKDGFADKLRQATSADQVIELFDQASEKTEELVQAPANDSGDFIVA VTACTTGIAHTYMAQEALQKVAAEMGVGIKVETNGASGVGNQLTAEDIRKAKAIIIAADKAVEMDRFDGKPLINRPVADGIRKTE ELINLALSGDTEVYRAANGAKAATASNEKQSLGGALYKHLMSGVSQMLPFVIGGGIMIALAFLIDGALGVPNENLGNLGSYHELA SMFMKIGGAAFGLMLPVFAGYVAYSIAEKPGLVAGFVAGAIAKEGFAFGKIPYAAGGEATSTLAGVSSGFLGALVGGFIAGALVL AIKKYVKVPRSLEGAKSILLLPLLGTILTGFVMLAVNIPMAAINTAMNDFLGGLGGGSAVLLGIVLGGMMAVDMGGPVNKAAYV FGTGTLAATVSSGGSVAMAAVMAGGMVPPLAIFVATLLFKDKFTKEERNSGLTNIIMGLSFITEGAIPFGAADPARAIPSFILGSAVA GGLVGLTGIKLMAPHGGIFVIALTSNALLYLVSVLVGAIVSGVVYGYLRKPQAZ

(SEO. ID. NO. 96)

MANKNTSTTRRPSKAELERKEAIQRMLISLGIAILLIFAAFKLGAAGITLYNLIRLLVGSLAYLAIFGLLIYLFFFKWIRKQEGLLSGF FTIFAGLLLIFEAYLVWKYGLDKSVLKGTMAQVVTDLTGFRTTSFAGGGLIGVALYIPTAFLFSNIGTYFIGSILILVGSLLVSPWSVY DIAEFFSRGFAKWWEGHERRKEERFVKQEEKARQKAEKEARLEQEETEKALLDLPPVDMETGEILTEEAVONLPPIPEEKWVEPEII LPOAELKFPEQEDDSDDEDVQVDFSAKEALEYKLPSLQLFAPDKPKDQSKEKKIVRENIKILEATFASFGIKVTVERAEIGPSVTKYE VKPAVGVRVNRISNLSDDLALALAAKDVRIEAPIPGKSLIGIEVPNSDIATVSFRELWEQSQTKAENFLEIPLGKAVNGTARAFDLSK MPHLLVAGSTGSGKSVAVNGIIASILMKARPDQVKFMMVDPKMVELSVYNDIPHLLIPVVTNPRKASKALQKVVDEMENRYELFA KVGVRNIAGFNAKVEEFNSQSEYKQIPLPFIVVIVDELADLMMVASKEVEDAIIRLGOKARAAGIHMILATORPSVDVISGLIKANV PSRVAFAVSSGTDSRTILDENGAEKLLGRGDMLFKPIDENHPVRLQGSFISDDDVERIVNFIKTQADADYDESFDPGEVSENEGEFS DGDAGGDPLFEEAKSLVIETQKASASMIQRRLSVGFNRATRLMEELEIAGVIGPAEGTKPRKVLOOZ

(SEQ. ID. NO. 97)

MSYFKKYKFDKSQFKLGMRTFKTGIAVFLVLLIFGFFGWKGLQIGALTAVFSLRESFDESVHFGTSRILGNSIGGLYALVFFLLNTFF HEAFWYTLVVVPICTMLTIMTNVAMNNKAGVIGGVAAMLIITLSIPSGETILYVFVRVLETFMGVFVAIIVNYDIDRIRLFLEKKEKZ

MNKSEHRHQLIRALITKNKIHTQAELQALLAENDIQVTQATLSRDIKNMNLSKVREEDSAYYVLNNGSISKWEKRLELYMEDALV WMRPVQHQVLLKTLPGLAQSFGSIIDTLSFPDAIATLCGNDVCLIICEDADTAQKCFEELKKFAPPFFFEEZ

(SEO. ID. NO. 99)

MKSIKLNALSYMGIRVLNIIFPILTGTYVARVLDRTDYGYFNSVDTILSFFLPFATYGVYNYGLRAISNVKDNKKDLNRTFSSLFYLC STLINYLISYFWIKRDIKLVKIHLSDFKPLFLPLTAMLVFANANMLFTFLDRLFLVKTGIDVNVSYYTIAORIVTVIAGVVTGAIGVSV PRLSYYLGKGDKEAYVSLVNRGSRIFNFFIIPLSFGLMVLGPNAILLYGSEKYIGGGILTSLFAFRTIILALDTILGSQILFTNGYEKRIT VYTVFAGLLNLGLNSLLFFNHIVAPEYYLLTTMLSETSLLVFYIIFIHRKQLIHLGHIFSYTVRYSLFSLSFVAIYFLINFVYPVDMVIN LPFLINTGLIVLLSAISYISLLVFTKDSIFYEFLNHVLALKNKFKKSZ

(SEO. ID. NO. 100)

MELFMKITNYEIYKLKKSGLTNQQILKVLEYGENVDQELLLGDIADISGCRNPAVFMERYFQIDDAHLSKEFQKFPSFSILDDCYPW DLSEIYDAPVLLFYKGNLDLLKFPKVAVVGSRACSKQGAKSVEKVIQGLENELVIVSGLAKGIDTAAHMAALQNGGKTIAVIGTGL DVFYPKANKRLQDYIGNDHLVLSEYGPGEQPLKFHFPARNRIIAGLCRGVIVAEAKMRSGSLITCERAMEEGRDVFAIPGSILDGLS DGCHHLIOEGAKLVTSGODVLAEFEFZ

(SEQ. ID. NO. 101)

MKQLTVEDAKQIELEILDYIDTLCKKHNINYIINYGTLIGAVRHEGFIPWDDDIDLSMPREDYORFINIFOKEKSKYKLLSLETDKNY FNNFIKITDSTTKIIDTRNTKTYESGIFIDIFPIDRFDDPKVIDTCYKLESFKLLSFSKHKNIVYKDSLLKDWIRTAFWLLLRPVSPRYFA NKIEKEIQKYSRENGQYMAFIPSKFKEKEVFPSGTFDKTIDLPFENLSLPAPEKFDTILTQFYGDYMTLPPEEKRFYSHEFHAYKLED

(SEQ. ID. NO. 102)
MIKINHLTITQNKDLRDLVSDLTMTIQDGEKVAIIGEEGNGKSTLLKILMGEALSDFTIKGNIQSDYQSLAYIPQKVPEDLKKKTLHD YFFLDSIDLDYSILYRLAEELHFDSNRFASDQEIGNLSGGEALKIQLIHELAKPFEILFLDEPSNDLDLETVDWLKGOIOKTROTVIFIS HDEDFLSETADTIVHLRLVKHRKEAETLVEHLDYDSYSEQRKANFAKQSQQAANNQRAYDKTMEKHRRVKQNVETALRATKDS TAGRILLAKKMKTVLSQEKRYEKAAQSMTQKPLEEEQIQLFFSDIQPLPASKVLVQLEKENLSIDDRVLVQKLQLTVRGQEKIGIIGP NGVGKSTLLAKLQRLLNDKREISLGFMPQDYHKKLQLDLSPIAYLSKTGEKEELQKIQSHLASLNFSYPEMQHQIRSLSGGQQGKL LLLDLVLRKPNFLLLDEPTRNFSPTSQPQIRKLFATYPGGLITVSHDRRFLKEVCSIIYRMTEHGLKLVNLEDLZ

(SEQ. ID. NO. 103)

MKPKTFYNLLAEQNLPLSDQQKEQFERYFELLVEWNEKINLTAITDKEEVYLKHFYDSIAPILQGLIPNETIKLLDIGAGAGFPSLPM KILYPELDVTIIDSLNKRINFLQLLAQELDLNGVHFYHGRAEDFAQDKNFRAQYDFVTARAVARMQVLSELTIPYLKVGGKLLALK ASNAPEELLEAKNALNLLFSKVEDNLSYALPNRDPRYITVVEKKKETPNKYPRKAGMPNKRPLZ

MSIKLIAVDIDGTLVNSQKEITPEVFSAIQDAKEAGVKVVIATGRPIAGVAKLLDDLOLRDEGDYVVTFNGALVOETATGHEIISESL TYEDYLDMEFLSRKLGVHMHAITKDGIYTANRNIGKYTVHESTLVSMPIFYRTPEEMAGKEIVKCMFIDEPEILDAAIEKIPAEFYER YSINKSAPFYLELLKKNVDKGSAITHLAEKLGLTKDETMAIGDEENDRAMLEVVGNPVVMENGNPEIKKIAKYITKTNDESGVAHAIRTWVLZ

(SEQ. ID. NO. 105)

MTWIILGVIALIVIFVIVSYNGLVKNRMQTKEAWSQIDVQLKRRNDLLPNLIETVKGYAKYEGSTLEKVAELRNQVAAATSPAEAM KASDALTRQVSGIFAVAESYPDLKASANFVKLQEELTNTENKISYSRQLYNSVVSNYNVKLETFPSNIIAGMFGFKAADFLQTPEEE KSVPKVDFSGLGDZ

(SEQ. ID. NO. 106)

MLFDQIASNKRKTWILLLVFFLLLALVGYAVGYLFIRSGLGGLVIALIIGFIYALSMIFQSTEIVMSMNGAREVDEQTAPDLYHVVE DMALVAQIPMPRVFIIDDPALNAFATGSNPQNAAVAATSGLLAIMNREELEAVMGHEVSHIRNYDIRISTIAVALASAITMLSSMAG RMMWWGGAGRRSDDDRDGNGLEIIMLVVSLLAIVLAPLAATLVQLAISRQREFLADASSVELTRNPQGMINALDKLDNSKPMS RHVDDASSALYINDPKKGGGFQKLFYTHPPISERIERLKQMZ

(SEQ. ID. NO. 107)

MKLNIQEIRKQSEGLNFEQTLDLVDDLRARNQEILDVKDILAVGKVQYEDRMYFLDYQLSYTIVLASSRSMEPVELVESYPVTEVF MEGATNQLDQEVLDDDLVLPIENGELDLAESVSDNILLNIPIKVLTAEEEAGQGFISGNDWQIMTEEEYQAQKAVKKEENSPFAGL OGLFDGDEZ

(SEQ. ID. NO. 108)

MKRQLALVVFSGGQDSTTCLFWVMQHYETVEAVTFAYGQRHHLEIQITREIAKEQGIRHHILDMSLLGQITAQPDFATIHISYIPDK LCVESKSLKLYLFSYRNHGDFHENCINTIGKDLVNLLDPRYLEVWGKFTPRGGISIDPYYNYGKQGTKYEGLAEQRLFQHDLYPEK IDNRZ

(SEQ. ID. NO. 109)

(SEQ. ID. NO. 110)

MGVETWFYSSICWLAIGLGSVWKFPYMTAANGGGGFLLIFLISTILIGFPLLLAEFALGRSAGVSAIKTFGKLGKNNKYNFIGWIGA FALFILLSFYSVIGGWILVYLGIEFGKLFQLGCTGDYAQLFTSIISNPAIALGAQAAFILLNIFIVSRGVQKGIERASKVMMPLLFIVFV FIIGRSLSLPNAMEGVLYFLKPDFSKLTSTGLLYALGQSFFALSLGVTVMLTYASYLDKKTNLVQSGISIVAMNISISIMAGLAIFQAR SPFNIQSEGGPSLLFIVLPQLFDKMPFGTIFYVLFLLLFLFATVTFSVVMLEINVDNITNQDNSKRAKWSVILGILTFVFGIPSALSYGV MADVHIFGKTFFDAMDFLVSNLLMPFGALYLSLFTGYIFKKALAMEELHLDERAWKQGLFQVWLFLLRFFVSSFQSSSLWSSLPNL CNQKGLEZ

(SEQ. ID. NO. 111)

MLKKWQLKDVILLAFLSIFFGGVFVGSGYVYNILSLLLTPLGLQAFANEILFGLWCMAAPIAAIFVPRVGSATIGEVLAALAEVLYG SQFGLGALLSGFVQGLGSEFGFIVTKNRYESWLSLTANSIGITLVSFVYEYIKLGYYAFSLPFVLSLLVVRFISVYFFCTILVRAIVKL YHQFATGGKAZ

(SEQ. ID. NO. 112)

MVKVATQTPIISLFLLILSLETSFIPSIALTLSVVAFCILFMLYYRRFKMLAWMIILAILPSFANYWAVQLHGDASQAVMLGTRAFVT VCIGLVFVSSVSLKELLLYLAQKGLSRSWSYALIVVFNSFPLIQQEIKSLKEACLLRGQELHFWSPLIYSKVLMTVFRWRHLYLRAL SAHGYDEHAQLKNSYRTFYIPKKTKLIYLLFFLLLQTSLFLZ

(SEQ. ID. NO. 113)

MRKHQLQVHKLTILSMMIALDVVLTPIFRIEGMAPMSSVVNILAGIMMGPVYALAMATVTAFIRMTTQGIPPLALTGATFGALLAG LFYKYGRKFHYSALGEILGTGIIGSIVSYPVMVLFTGSAAKLSWFIYTPRFFGATLIGTAISFIAFRFLIKQEFFKKVQGYFFSERIDZ

(SEQ. ID. NO. 114)

MQEFTNPFPIGSSSLIHCITNEISCEMLANGILALGCKPVMADDSREVLDFTKQSQALFINLGHLSAEKEKAIRMAASYANQSSLPMV VDAVGVTTSSIRKSLVKDLLDYRPTVLKGNMSEIRSLVGLKHHGVGVDASAKDQETEDLLQVLKDWCQTYPGMSFLVTGPKDLV VSKNQVAVLGNGCTELDWITGTGDLVGALTAVFLSQGKTGFEASCLAVSYLNIAAEKIVVQGMGLEEFRYQVLNQLSLLRRDEN WLDTIKGEVYEZ

(SEQ. ID. NO. 115)

 $MNHKIAILSDVHGNATALEAVIADAKNQGASEYWLLGDIFLPGPGANDLVALLKDLPITASVRGNWDDRVLEALDGQYGLEDPQ\\ EVQLLRMTQYLMERMDPATIVWLRSLPLLEKKEIDGLRFSISHNLPDKNYGGDLLVENDTEKFDQLLDAETDVAVYGHVHKQLL\\ RYGSQGQQIINPGSIGMPYFNWEALKNHRSQYAVIEVEDGELLNIQFRKVAYDYEAELELAKSKGLPFIEMYEELRRDDNYQGHNL\\ ELLASLIEKHGYVEDVKNFFDFLZ$

(SEQ. ID. NO. 116)

MNVNQIVRIIPTLKANNRKLNETFYIETLGMKALLEESAFLSLGDQTGLEKLVLEEAPSMRTRKVEGRKKLARLIVKVENPLEIEGIL SKTDSIHRLYKGQNGYAFEIFSPEDDLILIHAEDDIASLVEVGEKPEFQTDLASISLSKFEISMELHLPTDIESFLESSEIGASLDFIPAQG QDLTVDNTVTWDLSMLKFLVNELDIASLRQKFESTEYFIPKSEKFFLGKDRNNVELWFEEVZ

(SEQ. ID. NO. 117)

MKWTKIIKKIEEQIEAGIYPGASFAYFKDNQWTEFYLGQSDPEHGLQTEAGLVYDLASVSKVVGVGTVCTFLWEIGQLDIDRLVID FLPESDYPDITIRQLLTHATDLDPFIPNRDLLTAPELKEAMFHLNRRSQPAFLYSDVHFLLLGFILERIFNQDLDVILKDQVWKPWGM TETKFGPVELAVPTVRGVEAGIVHDPKARLLGRHAGSAGLFSTIKDLQIFLEHYLADDFARDLNQNFSPLDDKERSLAWNLEGDW LDHTGYTGTFIMWNRQKQEATIFLSNRTYEKDERAQWILDRNQVMNLIRKEEZ

(SEQ. ID. NO. 118)

MMKKTYNHILVWGVIFYSICIVCFCFTPQEQSTVGVGTPGIQHLGRLVFLLTPFNSLWKLGEVSDIGQLCWIFLQNILNVFLFFPLIFQ LLYLFPNLRKTKKVLLFSFLVSLGIECTQLILDFFFDFNRVFEIDDLWTNTLGGYLAWLLYKRLHKNKVRNZ

(SEQ. ID. NO. 119)

MKIPLLTFARHKFVYVLLTLLFLALVYRDVLMTYFFFDIHAPDLAKFDGQAIKNDLLKSALDFRILQFNLGFYQSFIIPIIIVLLGFQYI ELKNKVLRLSIGREVSYQGLKRKLTLQVASIPCLIYLVTVLIIAIITYFFGTFSPLGWNSLFSDGSGLQRLLDGEIKSYLFFTCVLLIGIF INAIYFLQIVDYVGNVTRSAITYLMFLWLGSMLLYSALPYYMVPMTSLMQASYGDVSLMKLFTPYILYIVPYMVLEKYEDNVZ

(SEQ. ID. NO. 120)

MFKVLQKVGKAFMLPIAILPAAGLLLGIGGALSNPTTIATYPILDNSIFQSIFQVMSSAGEVVFSNLSLLLCVGLCIGLAKRDKGTAA LAGVTGYLVMTATIKALVKLFMAEGSAIDTGVIGALVVGIVAVYLHNRYNNIQLPSALGFFGGSRFVPIVTSFSSILIGFVFFVIWPPF QQLLVSTGGYISQAGPIGTFLYGFLMRLSGAVGLHHIIYPMFWYTELGGVETVAGQTVVGAQKIFFAQLADLAHSGLFTEGTRFFA GRFSTMMFGLPAACLAMYHSVPKNRRKKYAGLFFGVALTSFITGTIEPIEFMFLFVSPVLYVVHAFLDGVSFFIADVLNISIGNTFSG GVIDFTLFGILQGNAKTNWVLQIPFGLIWSVLYYIIFRWFITQFNVLTPGRGEEVDSKEISESADSTSNTADYLKQDSLQIIRALGGSN NIEDVDACVTRLRVAVKEVNOVDKALLKQIGAVDVLEVKGGIQAIYGAKAILYKNSINEILGVDDZ

(SEO. ID. NO. 121)

MKFRKLACTVLAGAAVLGLAACGNSGGSKDAAKSGGDGAKTEITWWAFPVFTQEKTGDGVGTYEKSIIEAFEKANPDIKVKLETI DFKSGPEKITTAIEAGTAPDVLFDAPGRIIQYGKNGKLAELNDLFTDEFVKDVNNENIVQASKAGDKAYMYPISSAPFYMAMNKK MLEDAGVANLVKEGWTTDDFEKVLKALKDKGYTFGSLFSSGQGGDQGTRAFISNLYSGSVTDEKVSKYTTDDPKFVKGLEKATS WIKDNLINNGSQFDGGADIQNFANGQTSYTILWAPAQNGIQAKLLEASKVEVVEVPFPSDEGKPALEYLVNGFAVFNNKDDKKVA ASKKFIQFIADDKEWGPKDVVRTGAFPVRTSFGKLYEDKRMETISGWTQYYSPYYNTIDGFAEMRTLWFPMLQSVSNGDEKPADA LKAFTEKANETIKKAMKQZ

(SEQ. ID. NO. 122)

MQSTEKKPLTAFTVISTIILLLLTVLFIFPFYWILTGAFKSQPDTIVIPPQWFPKMPTMENFQQLMVQNPALQWMWNSVFISLVTMFL VCATSSLAGYVLAKKRFYGQRILFAIFIAAMALPKQVVLVPLVRIVNFMGIHDTLWAVILPLIGWPFGVFLMKQFSENIPTELLESA KIDGCGEIRTFWSVAFPIVKPGFAALAIFTFINTWNDYFMQLVMLTSRNNLTISLGVATMQAEMATNYGLIMAGAALAAVPIVTVF LVFQKSFTQGITMGAVKGZ

(SEQ. ID. NO. 123)

MKMFKNFNNILLNRKIVLLLRIVLMMILINHLLSTAVQKQDAVIFFKRELISIFSYNDYSEANLEIPKLLLNLSLFMVGWLSVILLES DLADHYHHLIRYQSSSFFDYTRKRLVVISKFFTQDLFVWFLGLLPLGIHFKTVALFFLLAQLMMLYLLLSYLIALISAGAGFSFFLYF LAFVGQEWMMDHIVTVYLVLLSLLVMLIVSRLEEKFKKGZ

(SEQ. ID. NO. 124)

MGKGEMGKGVIGLEFDSEVLVNKAPTLQLANGKTATFLTQYDSKTLLFAVDKEDIGQEIIGIAKGSIESMHNLPVNLAGARVPGGV NGSKAAVHEVPEFTGGVNGTEPAVHEIAEYKGSDSLVTLTTKKDYTYKAPLAQQALPETGNKESDLLASLGLTAFFLGLFTLGKK REOZ

(SEQ. ID. NO. 125)

MKKTFILVLGLFCLLPLSVFAIDFKINSYQGDLYIHADNTAEFRQKIVYQFEEDFKGQIVGLGRAGKMPSGFDIDPHPKIQAAKNG AELADVTSEVTEEADGYTVRVYNPGQEGDIVEVDLVWNLKNLLFLYDDIAELNWQPLTDSSESIEKFEFHVRGDKGAEKLFFHTG KLFREGTIEKSNLDYTIRLDNLPAKRGVELHAYWPRTDFASARDQGLKGNRLEEFNKIEDSIVREKDQSKQLVTWVLPSILSISLLLS VCFYFIYRRKTTPSVKYAKNHRLYEPPMELEPMVLSEAVYSTSLEEVSPLVKGAGKFTFDQLIQATLLDVIDRGNVSIISEGDAVGL RLVKEDGLSSFEKDCLNLAFSGKKEETLSNLFADYKVSDSLYRRAKVSDEKRIQARGLQLKSSFEEVLNQMQEGVRKRVSFWGLP DYYRPLTGGEKALQVGMGALTILPLFIGFGLFLYSLDVHGYLYLPLPILGFLGLVLSVFYYWKLRLDNRDGVLNEAGAEVYYLWT SFENMLREIARLDQAELESIVVWNRLLVYAATLFGYADKVSHLMKVHQIQVENPDINLYVAYGWHSTFYHSTAQMSHYASVANTA STYSVSSGSGSSGGGFSGGGGGGGGIGAFZ

(SEQ. ID. NO. 126)

MKKVRKIFQKAVAGLCCISQLTAFSSIVALAETPETSPAIGKVVIKETGEGGALLGDAVFELKNNTDGTTVSQRTEAQTGEAIFSNIK PGTYTLTEAQPPVGYKPSTKQWTVEVEKNGRTTVQGEQVENREEALSDQYPQTGTYPDVQTPYQIIKVDGSEKNGQHKALNPNPY ERVIPEGTLSKRIYQVNNLDDNQYGIELTVSGKTVYEQKDKSVPLDVVILLDNSNSMSNIRNKNARRAERAGEATRSLIDKITSDSE NRVALVTYASTIFDGTEFTVEKGVADKNGKRLNDSLFWNYDQTSFTTNTKDYSYLKLTNDKNDIVELKNKVPTEAEDHDGNRLM YQFGATFTQKALMKADEILTQQARQNSQKVIFHITDGVPTMSYPINFNHATFAPSYQNQLNAFFSKSPNKDGILLSDFITQATSGEH TIVRGDGQSYQMFTDKTVYEKGAPAAFPVKPEKYSEMKAAGYAVIGDPINGGYIWLNWRESILAYPFNSNTAKITNHGDPTRWYY NGNIAPDGYDVFTVGIGINGDPGTDEATATSFMQSISSKPENYTNVTDTTTKILEQLNRYFHTIVTEKKSIENGTITDPMGELIDLQLGT DGRFDPADYTLTANDGSRLENGQAVGGPQNDGGLLKNAKVLYDTTEKRIRVTGLYLGTDEKVTLTYNVRLNDEFVSNKFYDTNG TTTLHPKEVEQNTVRDFPIPKIRDVRKYPEITISKEKKLGDIEFIKVNKNDKKPLRGAVFSLQKQHPDYPDIYGAIDQNGTYQNVRT GEDGKLTFKNLSDGKYRLFENSEPAGYKPVQNKPIVAFQIVNGEVRDVTSIVPQDIPAGYEFTNDKHYITNEPIPPKREYPRTGGIG MLPFYLIGCMMMGGVLLYTRKHPZ



(SEQ. ID. NO. 127)

MKSINKFLTMLAALLLTASSLFSAATVFAAGTTTTSVTVHKLLATDGDMDKIANELETGNYAGNKVGVLPANAKEIAGVMFVWT NTNNEIIDENGQTLGVNIDPQTFKLSGAMPATAMKKLTEAEGAKFNTANLPAAKYKIYEIHSLSTYVGEDGATLTGSKAVPIEIELP LNDVVDAHVYPKNTEAKPKIDKDFKGKANPDTPRVDKDTPVNHQVGDVVEYEIVTKIPALANYATANWSDRMTEGLAFNKGTV KVTVDDVALEAGDYALTEVATGFDLKLTDAGLAKVNDQNAEKTVKITYSATLNDKAIVEVPESNDVTFNYGNNPDHGNTPKPNK PNENGDLTLTKTWVDATGAPIPAGAEATFDLVNAQTGKVVQTVTLTTDKNTVTVNGLDKNTEYKFVERSIKGYSADYQEITTAGE IAVKNWKDENPKPLDPTEPKVVTYGKKFVKVNDKDNRLAGAEFVIANADNAGQYLARKADKVSQEEKQLVVTTKDALDRAVA AYNALTAQQQTQQEKEKVDKAQAAYNAAVIAANNAFEWVADKDNENVVKLVSDAQGRFEITGLLAGTYYLEETKQPAGYALL TSRQKFEVTATSYSATGQGIEYTAGSGKDDATKVVNKKITIPOTGGIGTIIFAVAGAAIMGIAVYAYVKNNKDEDOLAZ

(SEO, ID, NO. 128)

MTMQKMQKMISRIFFVMALCFSLVWGAHAVQAQEDHTLVLQLENYQEVVSQLPSRDGHRLQVWKLDDSYSYDDRVQIVRDLHS WDENKLSSFKKTSFEMTFLENQIEVSHIPNGLYYVRSIIQTDAVSYPAEFLFEMTDQTVEPLVIVAKKTDTMTTKVKLIKVDQDHNR LEGVGFKLVSVARDVSEKEVPLIGEYRYSSSGQVGRTLYTDKNGEIFVTNLPLGNYRFKEVEPLAGYAVTTLDTDVQLVDHQLVTI TVVNQKLPRGNVDFMKVDGRTNTSLQGAMFKVMKEESGHYTPVLQNGKEVVVTSGKDGRFRVEGLEYGTYYLWELQAPTGYV OLTSPVSFTIGKDTRKELVTVVKNNKRPRIDVPDTGEETLVYLDACCHFVVWZ

(SEQ. ID. NO. 129)

MSHİYLSIFTSLLLMLGLVNVAQADEYLRIGMEAAYAPFNWTQDDDSNGAVKIDGTNQYANGYDVQIAKKIAKDLGKEPLVVKT KWEGLVPALTSGKIDMIIAGMSPTAERKQEIAFSSSYYTSEPVLLVKKDSAYASAKSLDDFNGAKITSQQGVYLYNLIAQIPGAKKE TAMGDFAQMRQALEAGVIDAYVSERPEALTAEAANSKFKMIQVEPGFKTGEEDTAIAIGLRKNDNRISQINASIETISKDDQVALM DRMIKEQPAEATTTEETSSSFFSQVAKILSENWQQLLRGAGITLLISIVGTIIGLIIGLAIGVFRTAPLSENKVIYGLQKLVGWVLNVYI EIFRGTPMIVQSMVIYYGTAQAFGINLDRTLAAIFIVSINTGAYMTEIVRGGILAVDKGQFEAATALGMTHNQTMRKIVLPQVVRNI LPATGNEFVINIKDTSVLNVISVVELYFSGNTVATQTYQYFQTFTIIAVIYFVLTFTVTRILRFIERRMDMDTYTTGANQMQTEDLKZ

(SEQ. ID. NO. 130)

MTQAILEIKHLKKSYGQNEVLKDISLTVHKGEVISIIGSSGSGKSTFLRSINLLETPTDGQILYHGQNVLEKGYDLTQYREKLGMVFQ SFNLFENLNVLENTIVAQTTVLKRERTEAEKIAKENLEKVGMGERYWQAKPKQLSGGQKQRVAIARALSMNPDAILFDEPTSALD PEMVGEVLKIMQDLAQEGLTMIVVTHEMEFARDVSHRVIFMDKGVIAEEGKPEDLFTNPKEDRTKEFLQRYLKZ

(SEQ. ID. NO. 131)

MKKYQLLFKISAVFSYLFFVFSLSQLTLIVQNYWQFSSQIGNLFWIQNILSLLFIGVMIVVLVKTGHGYLFRIPRKKWLWYSILTVLV LVFQISFNVQTAKHVQSTAEGWAVLIGYSGTNFAELGIYIALFFLVPLMEELIYRGLLQHAFFKHSRFGLDLLLPSILFALPHFSSLPS LLDIFVFATVGIIFAGLTRYTKSIYPSYAVHVINNIVATFPFLLTFLHRVLGZ

(SEQ. ID. NO. 132)

MNKKQWLGLGLVAVAAVGLAACGNRSSRNAASSSDVKTKAAIVTDTGGVDDKSFNQSAWEGLQAWGKEHNLSKDNGFTYFQS
TSEADYANNLQQAAGSYNLIFGVGFALNNAVKDAAKEHTDLNYVLIDDVIKDQKNVASVTFADNESGYLAGVAAAKTTKTKQV
GFVGGIESEVISRFEAGFKAGVASVDPSIKVQVDYAGSFGDAAKGKTIAAAQYAAGADIVYQVAGGTGAGVFAEAKSLNESRPEN
EKVWVIGVDRDQEAEGKYTSKDGKESNFVLVSTLKQVGTTVKDISNKAERGEFPGGQVIVYSLKDKGVDLAVTNLSEEGKKAVE
DAKAKILDGSVKVPEKZ

(SEQ. ID. NO. 133)

MSKKLQQISVPLISVFLGILLGAIVMWIFGYDAIWGYEELFYTAFGSLRGIGEIFRAMGPLVLIGLGFAVASRAGFFNVGLPGQALAG WILSGWFALSHPDMPRPLMILATIVIALIAGGIVGAIPGILRAYLGTSEVIVTIMMNYIVLYVGNAFIHAFPKDFMQSTDSTIRVGAN ATYQTPWLAELTGNSRMNIGIFFAIIAVAVIWFMLKKTTLGFEIRAVGLNPHASEYAGISAKRTIILSMIISGALAGLGGAVEGLGTF QNVYVQGSSLAIGFNGMAVSLLAANSPIGILFAAFLFGVLQVGAPGMNAAQVPSELVSIVTASIIFFVSVHYLIERFVKPKKQVKGG KZ

(SEQ. ID. NO. 134)

MGVKKKLKLTSLLGLSLLIMTACATNGVTSDITAESADFWSKLVYFFAEIIRFLSFDISIGVGIILFTVLIRTVLLPVFQVQMVASRKM QEAQPRIKALREQYPGRDMESRTKLEQEMRKVFKEMGVRQSDSLWPILIQMPVILALFQALSRVDFLKTGHFLWINLGSVDTTLVL PILAAVFTFLSTWLSNKALSERNGATTAMMYGIPVLIFIFAVYAPGGVALYWTVSNAYQVLQTYFLNNPFKIIAEREAVVQAQKDL ENRKRKAKKKAQKTKZ

(SEQ. ID. NO. 135)

 $MVIDPFAINELDYYLVSHFHSDHIDPYTAAAILNNPKLEHVKFIGPYHCGRIWEGWGVPKERIIVVKPGDTIELKDMKIHAVESFDR\\ TCLVTLPVNGADETGGELAGLAVTDEEMAQKAVNYIFETPGGTIYHGADSHFSNYFAKHGKDFKIDVALNNYGENPVGIQDKMT\\ SIDLLRMAENLRTKVIIPVHYDIWSNFMASTNEILELWKMRKDRLQYDFHPFIWEVGGKYTYPQDQHLVEYHHPRGFDDCFEQDS\\ NIQFKALLZ\\$

(SEO, ID, NO, 136)

MFLSGWLSSFANTYIHDLLGVLFPDSPFLNAFESAIAAPLVEEPLKLLSLVFVLALIPVRKLKSLFLLGIASGLGFQMIKDIGYIRTDLP EGFDFTISRILERIISGIASHWTFSGLAVVGVYLLYRAYKGQKVGKKQGLIFLGLALGTHFLFNSPFVELETELPLAIPVVTAIALYGF YHAYCFVEKHNELMTZ

(SEO, ID, NO, 137)

MKVEPRCDVLSRMSHFFIRILIMELQELVERSWAIRQAYHELEVKHHDSKWTVEEDLLALSNDIGNFQRLVMTKQGRYYDETPYT LEQKLSENIWWLLELSQRLDIDILTEMENFLSDKEKQLNVRTWKZ

(SEQ. ID. NO. 138)

MLDWKQFFLAYLRSRSRLFIYLLSLAFLVLLFQFLFASLGIYFLYFFFLCCFVTILFFTWDILVETQVYRQELLYGEREAKSPLEIALA EKLEAREMELYQQRSKAERKLTDLLDYYTLWVHQIKTPIAASQLLVAEVVDRQLKQQLEQEIFKIDSYTNLVLQYLRLESFHDDLV

LKQVQIEDLVKEIIRKYALFFIQKGLNVNLHDLDKEIVTDKKWLLVVIEQIISNSLKYTKEGGLEIYMDDQELCIKDTGIGIKNSDVLRVFERGFSGYNGRLTQQSSGLGLYLSKKISEELGHQIRIESEVGKGTTVRIQFAQVNLVLEZ

(SEQ. ID. NO. 139)

MELNTHNAEILLSAANKSHYPQDELPEIALAGRSNVGKSSFINTMLNRKNLARTSGKPGKTQLLNFFNIDDKMRFVDVPGYGYAR VSKKEREKWGCMIEEYLTTRENLRAVVSLVDLRHDPSADDVQMYEFLKYYEIPVIIVATKADKIPRGKWNKHESAIKKKLNFDPS DDFILFSSVSKAGMDEAWDAILEKLZ

(SEQ. ID. NO. 140)

MTKKQLHLVIVTGMSGAGKTVAIQSFEDLGYFTIDNMPPALLPKFLQLVEIKEDNPKLALVVDMRSRSFFSEIQAVLDELENQDGL DFKILFLDAADKELVARYKETRRSHPLAADGRILDGIKLERELLAPLKNMSQNVVDTTELTPRELRKTLAEQFSDQEQAQSFRIEV MSFGFKYGIPIDADLVFDVRFLPNPYYLPELRNQTGVDEPVYDYVMNHPESEDFYQHLLALIEPILPSYQKEGKSVLTIAMGCTGGQ HRSVAFAKRLAQDLSKNWSVNEGHRDKDRRKETVNRSZ

(SEQ. ID. NO. 141)

MRKPKITVIGGGTGSPVILKSLREKDVEIAAIVTVADDGGSSGELRKNMQQLTPPGDLRNVLVAMSDMPKFYEKVFQYRFSEDAG AFAGHPLGNLIIAGLSEMQGSTYNAMQLLSKFFHTTGKIYPSSDHPLTLHAVFQDGTEVAGESHIVDHRGIIDNVYVTNALNDDTPL ASRRVVQTILESDMIVLGPGSLFTSILPNIVIKEIGRALLETKAEIAYVCNIMTQRGETEHFTDSDHVEVLHRHLGRPFIDTVLVNIEK VPQEYMNSNRFDEYLVQVEHDFVGLCKQVSRVISSNFLRLENGGAFHDGDLIVDELMRI IQVKKZ

(SEQ. ID. NO. 142)

MKNLIKLLIIRLIVNLADSVFYIVALWHVSNNYSSSMFLGIFIAVNYLPDLLLIFFGPVIDRVNPQKILIISILVQLAVAVIFLLLLNQISF WVIMSLVFISVMASSISYVIEDVLIPQVVEYDKIVFANSLFSISYKVLDSIFNSFASFLQVAVGFILLVKIDIGIFLLALFILLLLKFRTSN ANIENFSFKYYKREVLQGTKFILNNKLLFKTSISLTLINFFYSFQTVVVPIFSIRYFDGPIFYGIFLTIAGLGGILGNMLAPIVIKYLKSN QIVGVFLFLNGSSWLVAIVIKDYTLSLILFFVCFMSKGVFNIIFNSLYQQIPPHQLLGRVNTTIDSIISFGMPIGSLVAGTLIDLNIELVLI AISIPYFLFSYIFYTDNGLKEFSIYZ

(SEQ. ID. NO. 143)

MMSNKNKEILIFAILYTVLFMFDGVKLLASLMPSAIANYLVYVVLALYGSFLFKDRLIQQWKEIRKTKRKFFFGVLTGWLFLILMT VVFEFVSEMLKQFVGLDGQGLNQSNIQSTFQEQPLLIAVFACVIGPLVEELFFRQVLLHYLQERLSGLLSIILVGLVFALTHMHSLAL SEWIGAVGYLGGGLAFSIIYVKEKENIYYPLLVHMLSNSLSLIILAISIVKZ

(SEO, ID, NO, 144)

LKKPIIEFKNVSKVFEDSNTKVLKDINFELEEGKFYTLLGASGSGKSTILNIIAGLLDATTGDIMLDGVRINDIPTNKRDVHTVFQSYA LFPHMNVFENVAFPLRLRKIDKKEIEQRVAEVLKMVQLEGYEKRSIRKLSGGQRQRVAIARAIINQPRVVLLDEPLSALDLKLRTD MQYELRELQQRLGITFVFVTHDQEEALAMSDWIFVMNDGEIVQSGTPVDIYDEPINHFVATFIGESNILPGTMIEDYLVEFNGKRFE AVDGGMKPNEPVEVVIRPEDLRITLPEEGKLQVKVDTQLFRGVHYEIIAYDELGNEWMIHSTRKAIVGEEIGLDFEPEDIHIMRLNE TEEEFDARIEEYVEIEQEAGLINAIEEERDEENKLZ

(SEQ. ID. NO. 145)

MKSMRILFLLALIQISLSSCFLWKECILSFKQSTAFFIGSMVFVSGICAGVNYLYTRKQEVHSVLASKKSVKLFYSMLLLINLLGAVL VLSDNLFIKNTLQQELVDFLLPSFFFLFGLDLLIFLPLKKYVRDFLAMLDRKKTVLVTILATLLFLRNPMTIVSLLIYIGLGLFFAAYL VPNSVKKEVSFYGHIFRDLVLVIVTLIFFZ

(SEQ. ID. NO. 146)

MVKKIIGMVLALLSVTVVGVGVFAYTIYQQGTETLAKTYKKIGEETKVIEATEPLTILLMGVDTGNVERTETWVGRSDSMILMTV NPKTKKTTMMSLERDILTRIESGNGQAHEAKLNSAYADGGAELAIETIQKMMNIHIDRYVMVNMRGLQKLVDAVGGITVNNILGF PISISDQEEFNTISIGVGEQHIGGEEALVYARMRYQDPEGDYGRQKRQREVIQKVMEKALSLNSIGHYQEILKALSDNMQTNIDLSA KSIPNLLGYKDSFKTIETQQLQGEGEILQGVSYQIVSRAHMLEMQNLLRRSLGQEEVTQLETNAVLFEDLFGRAPVGDEDNZ

(SEQ. ID. NO. 147)

MKKQAYVIIALTSFLFVFFFSHSLLEILDFDWSIFLHDVEKTEKFVFLLLVFSMSMTCLLALFWRGIEELSLRKMQANLKRLLAGQE VVQVADPDLDASFKSLSGKLNLLTEALQKAENQSLAQEEEIIEKERKRIARDLHDTVSQELFAAHMILSGISQQALKLDREKMQTQ LQSVTAILETAQKDLRVLLLHLRPVELEQKSLIEGIQILLKELEDKSDLRVSLKQNMTKLPKKIEEHIFRILQELISNTLRHAQASCLD VYLYQTDVELQLKVVDNGIGFQLGSLDDLSYGLRNIKERVEDMAGTVQLLTAPKQGLAVDIRIPLLDKEZ

(SEQ. ID. NO. 148)

MIVSIISQGFVWAILGLGIFMTFRILNFPDMTTEGSFPLGGAVAVTLITKGVNPFLATLVAVGAGCLAGMAAGLLYTKGKIPTLLSGI LVMTSCHSIMLLIMGRANLGLLGTKQIQDVLPFDSDLNQLLTGLIFVSIVIALMLFFLDTKLGQAYIATGDNPDMARSFGIHTGRME LMGLVLSNGVIALAGALIAQQEGYADVSRGIGVIVVGLASLIIGEVIFKSLSLAERLVTIVVGSIAYQFLVWAVIALGFNTSYLRLYS ALILAVCLMIPTFKQTILKGAKLSKZ

(SEQ. ID. NO. 149)

MKKMKVWSTVLATGVALTTLAACSGGSNSTTASSSEEKADKSQELVIYSNSVSNGRGDWLTAKAKEAGFNIKMVDIAGAQLADR VIAEKNNAVADMVFGIGAVDSNKIRDQKLLVQYKPKWLDKIDQSLSDKDNYYNPVIVQPLVLIGAPDVKEMPKDWTELGSKYKG KYSISGLQGGTGRAILASILVRYLDDKGELGVSEKGWEVAKEYLKNAYTLQKGESSIVKMLDKEDPIQYGMMWGSGALVGQKEQ NVVFKVMTPEIGVPFVTEQTMVLSTSKKQALAKEFIDWFGQSEIQVEYSKNFGSIPANKDALKDLPEDTKKFVDQVKPQNIDWEA VGKHLDEWVEKAELEYVQZ

(SEQ. ID. NO. 150)

MIKFDNIQIKYGDFVAIDNLNLDIHEGEFFTFLGPSGCGKSTTLRALVGFLDPSSGSIEVNGTDVTHLEPEKRGIGIVFQSYALFPTMT VFDNIAFGLKVKKVAPDVIKAKVSAVAAKIKISDQQLQRNVSELSGGQQQRVALARALVLEPKILCLDEPLSNLDAKLRVDLRKEL KRLQKELGITTLYVTHDQEEALTLSDRIAVFNNGYIEQVGTPVEIYHNSQTEFVCDFIGDINVLTDETVHEVLLKNTSVFLEDKKGYI RLEKVRFNRETEQDFILKGTIIDVEFSGVTIHYTIKVSESQILNVTSIDSQAAIRSVGESVELFITPSDVLOFZ

(SEQ. ID. NO. 151)
MRHKLNLKDWLIRLGLIWFLVTFIIYPNFDLVVNVFVKGGEFSLDAVHRVLKSQRALQSIMNSFKLAFSLIITVNVVGILCVLFTEYF
DIKGAKILKLGYMTSLIYGGVVLATGYKFVYGPYGLITKFLQNVIPSLDPNWFIGYGAVLFIMTFSGTANHTLFLTNTIRSVDYHTIE
AARNMGAKPFTVFRKVVLPTLIPTLFALTIMVFLSGLSAVAAPMIVGGKEFQTINPMIITFAGMGNSRDLAALLAIILGIATTILLTIM
NKIEKGGNYISISKTKAPLKKQKIASKPWNIIAHIVAYGLFTVFMLPLIFIVLYSFTDPVAIQTGNLTLSNFTLENYRLFFSNSAAFSPF
LVSFIYSIIAATTATILAVVFARVVRKHKSRFDFLFEYGALLPWLLPSTLLAVSLLFTFNQPQFLVLNQILVGSLVILLIAYIVVKIPFSY
RMVRAILFSVDDEMEDAARSMGASPFYTMMKVIIPFILPVVLSVIALNFNSLLTDFDLSVFLYHPLAQPLGITIRSAGDETATSNAQA LVFVYTIVLMIISGTVLYFTQRPGRKVRKZ



Table 3

ID201 - 4106.4 (SEO, ID, NO, 168)

AGCTATTTATTTGACCTTTCCTTTTTATTATTCAACTGGAGGGGGAAAAGTTTAATGAGAGCGCAAGAGTGTTTACGGAGT ATTTAAAGACTAAGACATCTGATGAAATTCCAAGCTTACTCCAGTCTTATTCAAAGTCCTTGACCATATCTGCTCACCTTAAA AGAGATATTGTAGATAAGCGGCTCCCTCTTGTGCATGACTTGGATATTAAAGATGGAAAGCTATCAAATTATATCGTGATGTT AGATATGTCTGTTAGTACAGCAGATGGTAAACAGGTAACCGTGCAATTTGTTCACGGGGTGGATGTCTACAAAGAAGCAAAGA ${\tt TTGCTCAATCCTCTTTTTTACATTTCAGAAGTGACTAGTAAAATGCAAGATTTGGATGACAATATTCGTTTTGATGAAAGTAG}$ $\tt GTAATGAGCAAATTGTAAAATTGCAAAATCAAAAGGTTTCCTTTGTCCGCGGAGCATCACATGAGTTGAAAACCCCTTTAGCC$ AGTCTTAGAATTATCCTAGAGAATATGCAGCATAATATTGGAGATTACAAAGATCATCCAAAATATATTGCAAAGAGTATAAA ${\tt TAAGATTGACCAGATGAGCCACTTATTAGAAGAAGTACTGGAGTCTTCTAAATTCCAAGAGTGGACAGAGTGTCGTGAGACCT}$ ${\tt TGACTGTTAAGCCAGTTTTAGTAGATATTTTATCACGTTATCAAGAATTAGCTCATTCAATAGGTGTTACAATTGAAAATCAA}$ TTGACAGATGCTACCAGGGTCGTCATGAGTCTTAGGGCATTGGATAAGGTTTTGACAAACCTGATTAGTAATGCAATTAAATA ${ t TCAGATAAAAATGGGCGTGTAATCATATCCGAGCAAGATGGCTATCTCTCTATCAAAAATACATGTGCGCCTCTAAGTGACC$ AAGAACTAGAACATTTATTTGATATATTCTATCATTCTCAAATCGTGACAGATAAGGATGAAAGTTCCGGTTTTGGGTCTTTAC ${ t ATTGTGAATAATATTTTAGAAAGCTATCAAATGGATTATAGTTTTCTCCCTTATGAACACGGTATGGAATTTAAGATTAGCTT$

(SEQ. ID. NO. 152)

MIKNPKLTKSFLRSFAILGGVGLVIHIAIYLTFPFYYIQLEGEKFNESARVFTEYLKTKTSDEIPSLLQSYSKSLTISAHLK RDIVDKRLPLVHDLDIKDGKLSNYIVMLDMSVSTADGKQVTVQFVHGVDVYKEAKNILLLYLPYTFLVTIAFSFVFSYFYTKR LLNPLFYISEVTSKMQDLDDNIRFDESRKDEVGEVGKQINGMYEHLLKVIYELESRNEQIVKLQNQKVSFVRGASHELKTPLA SLRIILENMQHNIGDYKDHPKYIAKSINKIDQMSHLLEEVLESSKFQEWTECRETLTVKPVLVDILSRYQELAHSIGVTIENQ LTDATRVVMSLRALDKVLTNLISNAIKYSDKNGRVIISEQDGYLSIKNTCAPLSDQELEHLFDIFYHSQIVTDKDESSGLGLY IVNNILESYQMDYSFLPYEHGMEFKISLZ

ID202 - 4106.9 (SEQ. ID. NO. 169)

(SEQ. ID. NO. 153)

MDKIIKTISESGAFRAFVLDSTETVRTAQEKHQTQASSTVALGRTLIASQILAANEKGNTKLTVKVLGSSSLGAIITVADTKG NVKGYVQNPGVDIKKTATGEVLVGPFVGNGQFLVITDYGTGNPYNSITPLISGEIGEDLAFYLTESQQTPSAVGLNVLLDEED KVKVAGGFLVQVLPGAKKEEIARFEKRIQEMPAISTLLESDDHIEALLKAIYGDEAYKRLSEEEIRFQCDCSHERFMNALASL PSSDLQEMKEEDHGAEITCQFCQTTYNFDEKDLEELIRDKSZ

ID203 - 4115 (SEQ. ID. NO. 170)

ATGAAATCAATAACTAAAAAGATTAAAGCAACTCTTGCAGGAGTAGCTGCCTTGTTTGCAGTATTTGCTCCATCATCTTTGTATC
TGCTCAAGAATCATCAACTACACTGTTAAAGAAGGTGATACACTTTCAGAAATCGCTGAAACTCACAACACAACAGTTGAAA
AATTGGCAGAAAAACAACCACATTGATAACATTCATTTGATTTATGTTTGATCAAGAAGTTGGTTATCGATGGCCCTGTAGCGCCT
GTTGCAACACCAGCGCCAGCTACTTATGCGGCACCAGCCGCTCAAGATGAAACTGTTTCAGCTCCAGTAGCAGAAACTCCAGT
AGTAAGTGAAACAGTTGTTTCAACTGTAAGCGGATCTGAAGCAGAAGCCCAAAGAATGGATCGCTCAAAAAGAATCAGGTGGTA
GTATACAGCTACAAATGGACGTTATATCGGACGTTACCAATTAA

(SEQ. ID. NO. 154)

MKSITKKIKATLAGVAALFAVFAPSFVSAQESSTYTVKEGDTLSEIAETHNTTVEKLAENNHIDNIHLIYVDQELVIDGPVAP VATPAPATYAAPAAQDETVSAPVAETPVVSETVVSTVSGSEAEAKEWIAQKESGGSIQLQMDVISDVTNZ

ID204 - 4117.1 (SEQ. ID. NO. 171)



(SEQ. ID. NO. 155)

MNLGEFWYNKINKNRGRRLMKKVRFIFLALLFFLASPEGAMASDGTWQGKQYLKEDGSQAANEWVFDTHYQSWFYIKADANYA ENEWLKQGDDYFYLKSGGYMAKSEWVEDKGAFYYLDQDGKMKRNAWVGTSYVGATGAKVIEDWVYDSQYDAWFYIKADGQHAE KEWLQIKGKDYYFKSGGYLLTSQWINQAYVNASGAKVQQGWLFDKQYQSWFYIKENGNYADKEWIFENGHYYYLKSGGYMAAN EWIWDKESWFYLKFDGKMAEKEWVYDSHSQAWYYFKSGGYMTANEWIWDKESWFYLKSDGKIAEKEWVYDSHSQAWYYFKSGGYMTANEWIWDKESWFYLKSDGKTTNENAAYYQVVPVTA NVYDSDGEKLSYISQGSVVLDKDRKRKEWVYDSHSQAWYYFKSGGYMAKNETVDGYQLGSDGKWLGGKTTNENAAYYQVVPVTA NVYDSDGEKLSYISQGSVVLDKDRKRKSDDKRLAITISGLSGYMKTEDLQALDASKDFIPYYESDGHRFYHYVAQNASIPVASH LSDMEVGKKYYSADGLHFDGFKLENPFLFKDLTEATNYSAEELDKVFSLLNINNSLLENKGATFKEAEEHYHINALYLLAHSA LESNWGRSKIAKDKNNFFGITAYDTTPYLSAKTFDDVDKGILGATKWIKENYIDRGRTFLGNKASGMNVEYASDPYWGEKIAS VMMKINEKLGGKDZ

ID205 - 4118.1 (SEQ. ID. NO. 172)

(SEQ. ID. NO. 156)

MKKLGTLLVLFLSAIILVACASGKKDTTSGQKLKVVATNSIIADITKNIAGDKIDLHSIVPIGQDPHEYEPLPEDVKKTSEAN LIFYNGINLETGGNAWFTKLVENAKKTENKDYFAVSDGVDVIYLEGQNEKGKEDPHAWLNLENGIIFAKNIAKQLSAKDPNNK EFYEKNLKEYTDKLDKLDKESKDKFNKIPAEKKLIVTSEGAFKYFSKAYGVPSAYIWEINTEEEGTPEQIKTLVEKLRQTKVP SLFVESSVDDRPMKTVSQDTNIPIYAQIFTDSIAEQGKEGDSYYSMMKYNLDKIAEGLAKZ

ID206 - 4119.1 (SEQ. ID. NO. 173)

(SEQ. ID. NO. 157)

MEWYKKIGLLATTGLALFGLGACSNYGKSADGTVTIEYFNQKKEMTKTLEEITRDFEKENPKIKVKVVNVPNAGEVLKTRVLA GDVPDVVNIYPQSIELQEWAKAGVFEDLSNKDYLKRVKNGYAEKYAVNEKVYNVPFTANAYGIYYNKDKFEELGLKVPETWDE FEQLVKDIVAKGQTPFGIAGADAWTLNGYNQLAFATATGGGKEANQYLRYSQPNAIKLSDPIMKDDIKVMDILRINGSKQKNW EGAGYTDVIGAFARGDVLMTPNGSWAITAINEQKPNFKIGTFMIPGKEKGQSLTVGAGDLAWSISATTKHPKEANAFVEYMTR PEVMQKYYDVDGSPTAIEGVKQAGEDSPLAGMTEYAFTDRHLVWLQQYWTSEADFHTLTMNYVLTGDKQGMVNDLNAFFNPMK ADVDZ

ID207 - 4123.1 (SEQ. ID. NO. 174)



(SEQ. ID. NO. 158)

MKKIKPHGPLPSQTQLAYLGDELAAFIHFGPNTFYDQEWGTGQEDPERFNPSQLDAREWVRVLKETGFKKLILVVKHHDGFVL YPTAHTDYSVKVSPWRRGKGDLLLEVSQAATEFDMDMGVYLSPWDAHSPLYHVDREADYNAYYLAQLKEILSNPNYGNAGKFA EVWMDGARGEGAQKVNYEFEKWFETIRDLQGDCLIFSTEGTSIRWIGNERGYAGDPLWOKVNPDKLGTEAELNYLOHGDPSGT IFSIGEADVSIRPGWFYHEDQDPKSLEELVEIYFHSVGRGTPLLLNIPPNQAGLFDAKDIERLYEFATYRNELYKEDLALGAE VSGPALSADFACRHLTDGLETSSWASDADLP1QLELDLGSPKTFDV1ELREDLKLGQR1AAFHVQVEVDGVWQEFGSGHTVGY KRLLRGAVVEAQKIRVVITESQALPLLTKISLYKTPGLSKKEVVQELAFAEKSLAVAKGENAYFTVKRRECSGPLEAKISIQP GTGVHGVAYQDEIQVLAFQTGETEKSLTLPTLYFAGDKTLDFYLNLTVDGQLVDQLQVQVSZ

ID208 - 4125.12 (SEQ. ID. NO. 175)

TGGGTGGCTTTCTGCCTGGCATTTATTGATTGATATTCTATTTGTAGTGGCATATTTGGGTGTTTTAACAACTAAGAGCCAGC GCCTATCTTGGCTATATTGGGGCCTCATGCTGACTTATGTAGTTGGGAATACTGCCTTTGTTGCTGTTAATTATATCTGGTTT ATAAATCTATTGCTTGCTGAAAATGAACGTAGTCGTATCGGTCAGGATTTGCATGATAGTCTGGGACATACCTTTGCTATGCT GAGTGTCAAGACAGATTTAGCCTTGCAGTTATTTCAGATGGAGGCTTATCCACAGGTGGAAAAGGAATTAAAAGAAATTCACC AGATCAGCAAGGATCCATGA

(SEO. ID. NO. 159)

MLERLKRIHYMFWISLIFMIFPILSVVTGWLSAWHLLIDILFVVAYLGVLTTKSQRLSWLYWGLMLTYVVGNTAFVAVNYIWF FFFLSNLLSYHFSVRSLKSLHVWTFLLAQVLVVGQLLIFQRIEVEFLFYLLVILTFVDLMTFGLVRIRIVEDLKEAQVKQNAQ INLLLAENERSRIGODLHDSLGHTFAMLSVKTDLALOLFOMEAYPOVEKELKEIHOISKDPZ

ID209 - 4126.3 (SEQ. ID. NO. 176)
ATGAATGATAAGTTAAAAATCTTCTTGTTGCTAGGAGTATTTTTTCTAGCCATAACCGGTTTCTATGTTCTATTGATACGAAA TGCAGGGCAGACAGATGCCTCGCAAATTGAAAAGGCGGCAGTTAGCCAAGGAGGAAAAAGCAGTGAAAAAAACAGAAATTAGTA AAGACGCAGACTTGCACGAAATTTATCTAGCTGGAGGTTGTTTCTGGGGAGTGGAGGAATATTTCTCACGTGTTCCCGGGGTG ACGGATGCCGTTTCAGGCTATGCAAATGGTAGAGGAGAAACAACCAAGTACGAATTGATTAACCAAACAGGTCATGCAGAAAC $\tt CGTCCATGTCACCTATGATGCCAAGCAAATTTCTCTCAAGGAAATCCTGCTTCACTATTTCCGCATTATCAATCCAACCAGCA$ AAAATAAACAAGGAAATGATGTGGGGACCCAGTACCGTACTGGTGTTTATTACACAGATGACAAGGATTTGGAAGTGATTAAC CAAGTCTTTGATGAGGTGGCTAAGAAATACGATCAACCTCTAGCAGTTGAAAAGGAAAACTTGAAGAATTTTGTGGTGGCTGA GGATTACCATCAAGACTATCTCAAGAAAAATCCAAATGGCTACTGCCATATCAATGTTAATCAGGCGGCCTATCCTGTCATTG ATGCCAGCAAATATCCAAAACCAAGTGATGAGGAATTGAAAAAGACCCTGTCACCTGAGGAGTATGCAGTTACCCAGGAAAAT ${\tt CAAACAGAACGAGCTTTCTCAAACCGTTACTGGGATAAATTTGAATCCGGTATCTATGTGGATATAGCAACTGGGGAACCTCT}$ AGGAAGATAAGTCCTACAATATGACGCGTATGGAAGTGCGGAGCCGAGTAGGAGATTCTCACCTTGGGCATGTCTTTACGGAT GGTCCACAGGACAAGGGCGGCTTACGTTACTGTATCAATAGCCTCTCTATCCGCTTTATTCCCAAAGACCAAATGGAAGAAAA AGGCTACGCTTATTTACTAGATTATGTTGATTAA

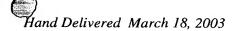
(SEQ. ID. NO. 160)

MNDKLKIFLLLGVFFLAITGFYVLLIRNAGQTDASQIEKAAVSQGGKAVKKTEISKDADLHEIYLAGGCFWGVEEYFSRVPGV TDAVSGYANGRGETTKYELINQTGHAETVHVTYDAKQISLKEILLHYFRIINPTSKNKQGNDVGTQYRTGVYYTDDKDLEVIN QVFDEVAKKYDQPLAVEKENLKNFVVAEDYHQDYLKKNPNGYCHINVNQAAYPVIDASKYPKPSDEELKKTLSPEEYAVTQEN QTERAFSNRYWDKFESGIYVDIATGEPLFSSKDKFESGCGWPSFTQPISPDVVTYKEDKSYNMTRMEVRSRVGDSHLGHVFTD GPQDKGGLRYCINSLSIRFIPKDQMEEKGYAYLLDYVDZ

ID210 - 4127.1 (SEQ. ID. NO. 177)

 ${ t ATGAAAAAGAAATGGATGTATTATGCTGCTTGTTCTTCTAATGAATCTGCCGATGACAGTTCATCTGATAAAGGAGACGGCGG}$ TTCGCTAGTCGTTTATTCACCAAACTCAGAGGGCTTAATTGGAGCAACTATTCCTGCCTTTGAAGAAAATATGGTATCAAAG TAGAACTGATTCAAGCTGGTACTGGAGAACTTTTCAAAAAACTAGAGTCAGAAAAAGAAGTTCCTGTAGCTGATGTTATCTTT GGTGGTTCTTATACACAATATACTACCCACGGAGAACTCTTTGAAAACTATACTTCAAAAGAAAATGATAATGTTATCAAAGA ATATCAAAACACAACTGGCTACTCTACTCCTTATACACTAGATGGTAGTGTTTTAATCGTCAACCCTGATTTAACTAAAGGCA TGAACATCGAAGGATATAACGATCTTTTCAAACCTGAACTAAAAGGAAAAATCGCAACTGCTGACCCAGCAAACTCTTCTAGC GCCTTTGCTCAATTAACAAATATGCTACAAGCTCAAGGTGGTTACAAAGATGATAAGGCTTGGTCTTATGTAAAAGATCTTTT CACACTTATTGATGGTAAAATCGGTTCAAGTTCATCTAGTGTCTATAAAGTAGTCGCTGATGGAGAAATGGCTGTTGGTCTCT CTTATGAAGATCCAGCAGTTAAACTCTTAAATGACGGAGCTAACATTAAGGTAGTCTATCCAAAAGAAGGAACCGTCTTCCTA CCTGCTAGTGCTGCTATCGTTAAAAAATCTAAAAATATGGAAAATGCCAAGAAATTTATCGATTTTATCTCTCAAGAAGT





ACAAGATACACTTGGTACAACCACTACTAACCGTCCTGTTCGTAAAAATGCTAAAACAAGCGAAAACATGAAACCAATTGACA AAATCAAAACACTCACTGAAGATTATGATTATGTCATCAAGAATAAATCAGATATCGTTAAGAAATACAACGAAGTCTTTACA GATATCCAATCTAAACAGTAA

(SEQ. ID. NO. 161)

MKKKWMYYAACSSNESADDSSSDKGDGGSLVVYSPNSEGLIGATIPAFEEKYGIKVELIQAGTGELFKKLESEKEVPVADVIFGGSYTQYTTHGELFENYTSKENDNVIKEYQNTTGYSTPYTLDGSVLIVNPDLTKGMNIEGYNDLFKPELKGKIATADPANSSSAFAQLTNMLQAQGGYKDDKAWSYVKDLFTLIDGKIGSSSSSVYKVVADGEMAVGLSYEDPAVKLLNDGANIKVVYPKEGTVFLPASAAIVKKSKNMENAKKFIDFIISQEVQDTLGTTTTNRPVRKNAKTSENMKPIDKIKTLTEDYDYVIKNKSDIVKKYNEVFTDIQSKQZ

ID211 - 4127.2 (SEO. ID. NO. 178)

(SEO. ID. NO. 162)

MSEIKIINAKKIYHDVPVIENLNITIPKGSLFTLLGASGCGKTTLLRMIAGFNSIEGGEFYFDDTKINNMEPSKRNIGMVFQN
YAIFPHLTVRDNVAFGLMQKKVPKEELIQQTNKYLELMQIAQYADRKPDKLSGGQQQRVTLACALAVNPSVLLMDEPLSNLEA
KLRLDMRQAIREIQHEVGITTVYVTHDQEEAMAISDQIAVMKDGVIQQIGRPKELYHKPANEFVATFIGRTNIIPANLEKRSD
GAYIVFSDGYALRMPALDQVEEQAIHVSIRPEEFIKDESGDIEGTIRDSVYLGLNTDYFIETGFASKIQVSEESTFEEDLQKG
NRIRLRINTQKLNIFSADGSONLIKGVNHGTZ

ID212 - 4136.1 (SEQ. ID. NO. 179)

(SEQ. ID. NO. 163)

MKKKLLAGAITLLSVATLAACSKGSEGADLISMKGDVITEHQFYEQVKSNPSAQQVLLNMTIQKVFEKQYGSELDDKEVDDTI AEEKKQYGENYQRVLSQAGMTLETRKAQIRTSKLVELAVKKVAEAELTDEAYKKAFDEYTPDVTAQIIRLNNEDKAKEVLEKA KAEGADFAQLAKDNSTDEKTKENGGEITFDSASTEVPEQVKKAAFALDVDGVSDVITATGTQAYSSQYYIVKLTKKTEKSSNI DDYKEKLKTVILTQKQNDSTFVQSIIGKELQAANIKVKDQAFQNIFTQYIGGGDSSSSSSTSNEZ

ID213 - 4137.3 (SEQ. ID. NO. 180)

ATGAAAAAAATATTAAACAATATGTAACCTTAGGTACTGTAGTGGTATTATCAGCATTTGTTGCTAACTCAGTTGCAGCTCA GGAGACTGAAACTTCTGAAGTATCAACACCAAAGTTGGTGCAACCTGTTGCACCAACGACTCCGATTTCGGAAGTACAACCTA ${\tt CATCGGATAACTCTTCGGAAGTTACTGTACAACCTCGAACAGTTGAAACTACTGTTAAGGATCCATCTTCTACAGCGGAAGAA}$ $\tt CGGTGACTTTACTGTAGTGATTAAGTACAATCAGTCAAGTGAGAAAGGCTTACAAGCTCTGTTTGGAATATCTAATTCCAAAC$ ${\tt CCGGTCAACAAAATAGTTATGTAGATGTTTCCTTAGAGACAATGGTGAGTTGGGGATGGAAGCGCGTGATACTTCTTCCAAT$ AGGATATTAAAGGTATTGATTACTATATGCTTGGGGGAGTGAAACGTGCAGGAAAAACGGCGTTTGGTTTTAACGGAACACTA GAAAATATCAAATTCTTTAATAGTGCATTGGATGAAGAAACTGTTAAAAAGATGACAAACGCTGTTACTGGACATTTAAT GCATTGACGCTCGTTACGGTGGAACTCATGATTTCTTGAATAAAATTAATATTGCTACAAGTTATAGTGATGATAATGGTAAG ACATGGACTAAACCAAAATTAACATTGGCATTCGATGATTTTGCGCCAGTACCATTAGAATGGCCTCGTGAAGTTGGTGGACG ATGTGATGCCTGCTGGAGTAAGTTTTAGAGAAGCAACTAGAAAAGATTCAGGTTATAAACAAATTGATGGTAATTATTACCTT AAATTAAGGAAACAAGGTGATACTGATTACAATTATACTATTCGTGAGAATGGTACTGTATACGACGATCGTACCAACAGACC AACTGAATTTTCAGTAGATAAAAATTTCGGTATTAAACAAAATGGTAATTATTTGACGGTAGAGCGG

(SEQ. ID. NO. 164)





MKKNIKQYVTLGTVVVLSAFVANSVAAQETETSEVSTPKLVQPVAPTTPISEVQPTSDNSSEVTVQPRTVETTVKDPSSTAEE TPVLEKNNVTLTGGGENVTKELKDKFTSGDFTVVIKYNQSSEKGLQALFGISNSKPGQQNSYVDVFLRDNGELGMEARDTSSN KNNLVSRPASVWGKYKQEAVTNTVAVVADSVKKTYSLYANGTKVVEKKVDNFLNIKDIKGIDYYMLGGVKRAGKTAFGFNGTL ENIKFFNSALDEETVKKMTTNAVTGHLIYTANDTTGSNYFRIPVLYTFSNGRVFSSIDARYGGTHDFLNKINIATSYSDDNGK TWTKPKLTLAFDDFAPVPLEWPREVGGRDLQISGGATYIDSVIVEKKNKQVLMFADVMPAGVSFREATRKDSGYKQIDGNYYL KLRKQGDTDYNYTIRENGTVYDDRTNRPTEFSVDKNFGIKQNGNYLTVER

ID214 - 4185 (SEQ. ID. NO. 181)

ATGAAAAAATTTAGCCTATTACTAGCTATCCTACCATTTTTGGTTGCCTGTGAGAATCAAGCTACACCCAAAGAGACTAGCGC
TCAAAAGACAATCGTCCTTGCTACAGCTGCGACGTGCCACCATTTTGACTACGAAGACAAGGGCAATCTGACAGGCTTTGATA
TCGAAGTTTTAAAGGCAGTAGATGAAAAAACTCAGCGACTACGAGATTCAATTCCAAAGAACCGCCTGGGAGAGACATCTTCCCA
GGACTTGATTCTGGTCACTATCAGGCTGCGGCCCAATAACTTGAGTTACACAAAAGAGCGTGCTGAAAAAATACCTTTACTCGCT
TCCAATTTCCAACAATCCCCTCGTCCTTGTCAGCAACAAGAAAAAATCCTTTGACTTCTCTTGACCAGATCGCTGGTAAAACAA
CACAAGAGGATACCGGAACTTCTAACGCTCAATTCATCAATAACTGGAATCAGAAACACACTGATAATCCCGCTACAATTAAT
TTTTCTGGTGAGGATATTGGTAAACGAATCTTAGACATTGCTTAACGGAGTTTTCATTTTTTAGACAAGGTATCCGT
TCAAAAAGATTATCAAGGACCGTGGTTTAGACCTCTCAGTCGTTGATTTACCTTCTGCAGATAGCCCCAGCAATTATATCATTT
TCTCAAGCGACCAAAAAGAGTTTAAAAGAGCAATTTGATAAAGCGCTCCAAAGAACCTCTATCAAGACGGAACCCTTGAAAAACTC
AGCAATACCTATCTAGGTGGTTCTTACCTCCCAGATCAATCTCAGTTTACAATAA

(SEO. ID. NO. 165)

MKKFSLLLAILPFLVACENQATPKETSAQKTIVLATAGDVPPFDYEDKGNLTGFDIEVLKAVDEKLSDYEIQFQRTAWESIFP GLDSGHYQAAANNLSYTKERAEKYLYSLPISNNPLVLVSNKKNPLTSLDQIAGKTTQEDTGTSNAQFINNWNQKHTDNPATIN FSGEDIGKRILDLANGEFDFLVFDKVSVQKIIKDRGLDLSVVDLPSADSPSNYIIFSSDQKEFKEQFDKALKELYQDGTLEKL SNTYLGGSYLPDQSQLQZ

ID215 - 4211.1 (SEQ. ID. NO. 182)

(SEQ. ID. NO. 166)

MKKNSLYIISSLFFACVLFVYATATNFQNSTSARQVKTETYTNTVTNVPIDIRYNSDKYFISGFASEVSVVLTGANRLSLASE MQESTRKFKVTADLTDAGVGTIEVPLSIEDLPNGLTAVATPQKITVKIGKKAQKDKVKIVPEIDPSQIDSRVQIENVMVSDKE VSITSDQETLDRIDKIIAVLPTSERITGNYSGSVPLQAIDRNGVVLPAVITPFDTIMKVTTKPVAPSSSTSNSSTSSSSETSS STKATSSKTNZ

ID216 - 4127.3 (SEQ. ID. NO. 183)

(SEO. ID. NO. 167)

MLIGEGYRTFPVLIYTQFISEVGGNSAFAIMAIIIALAIFLIQKHIANRYSFSMNLLHPIEPKKTTKGKMAAIYATVYGIIFI SVLPQIYLIYTSFLKTSGMVSVKGYSPNSYKVAFHRMGSAIFNTIRIPLIALVLVVLFATFISYLAVRKRNLFTNLIDSLSMV PYIVPGTVLGIAFISSFNTGLFGSGFLMITGTAFILIMSLSARRLPYTIRSSVASLQQIAPSIEEAAESLGSSRLNTFAKITT PMMLSGIISGAILSZ

J



Table 4

ID301 (SEQ. ID. NO. 196)

ATGAATAAGAAAAAATGATTTTAACAAGTCTAGCCAGCGTCGCTATCTTAGGGGGCTGGTTTTGTTACGTCTCAGCCTACTTT TGTAAGAGCAGAAGAATCTCCACAAGTTGTCGAAAAATCTTCATTAGAGAAAAAATATGAGGAAGCAAAAGCAAAAGCTGATA $\tt CTGCCAAGAAAGATTACGAAACGGCTAAAAAGAAAGCAGAAGACGCTCAGAAAAAGTATGAAGATGATCAGAAGAGAACTGAG$ GAGAAAGCTCGAAAAGAAGCAGAAGCATCTCAAAAATTGAATGTGGCGCTTGTTGTTCAAAATGCATATAAAGAGTACCG AGAAGTTCAAAATCAACGTAGTAAATATAAATCTGACGCTGAATATCAGAAAAAAATTAACAGAGGTCGACTCTAAAATAGAGA AGGCTAGGAAAGAGCAACAGGACTTGCAAAATAAATTTAATGAAGTAAGAGCAGTTGTAGTTCCTGAACCAAATGCGTTGGCT GAGACTAAGAAAAAAGCAGAAGAAGCTAAAGCAGAAGAAAAAGTAGCTAAGAGAAAATATGATTATGCAACTCTAAAGGTAGC ACTAGCGAAGAAGAAGTAGAGGCTAAGGAACTTGAAATTGAAAAACTTCAATATGAAATTTCTACTTTGGAACAAGAAGTTG CTACTGCTCAACATCAAGTAGATAATTTGAAAAAACTTCTTGCTGGTGCGGATCCTGATGATGGCACAGAAGTTATAGAAGCT CAGCCTTGATCCTGAAGGTAAGACTCAGGATGAATTAGATAAAGAAGCAGAAGAAGCTGAGTTGGATAAAAAAGCTGATGAAC $\tt TTCAAAATAAAGTTGCTGATTTAGAAAAAGAAATTAGTAACCTTGAAATATTACTTGGAGGGGCTGATCCTGAAGATGATACT$ TGATCCTGAAGGTAAGACTCAGGATGAATTAGATAAAGAAGCAGAAGAAGCTGAGTTGGATAAAAAAGCTGATGAACTTCAAA ATAAAGTTGCTGATTTAGAAAAAGAAATTAGTAACCTTGAAATATTACTTGGAGGGGCTGATTCTGAAGATGATACTGCTGCT $\tt CTTCAAAATTAAATTAGCTACTAAAAAAGCTGAATTGGAAAAAAACTCAAAAAGAATTAGATGCAGCTCTTAATGAGTTAGGCCC$ TGATGGAGATGAAGAAGAAACTCCAGCGCCGGCTCCTCAACCAGAGCAACCAGCTCCTGCACCAAAACCAGAGCAACCAGCTC CAGCTCCAAAACCAGAGCAACCAGCTCCTGCACCAAAACCAGAGCAACCAGCTCCAGCTCCAAAACCAGAGCAACCAGCTCCA GCTCCAAAACCAGAGCAACCAGCTAAGCCGGAGAAACCAGCTGAAGAGCCTACTCAACCAGAAAAACCAGCCACTCCAAAAAC AGGCTGGAAACAAGAAAACGGTATGTGGTATTTCTACAATACTGATGGTTCAATGGCAATAGGTTGGCTCCAAAACAACGGTT ${\tt CATGGTACTACCTAAACGCTAACGGCGCTATGGCAACAGGTTGGGTGAAAGATGGAGATACCTGGTACTATCTTGAAGCATCA}$ GGTGCTATGAAAGCAAGCCAATGGTTCAAAGTATCAGATAAATGGTACTATGTCAACAGCAATGGCGCTATGGCGACAGGCTG GCTCCAATACAATGGCTCATGGTACTCAACGCTAATGGTGATATGGCGACAGGATGGCTCCAATACAACGGTTCATGGT ATTACCTCAACGCTAATGGTGATATGGCGACAGGATGGGCTAAAGTCAACGGTTCATGGTACTACCTAAACGCTAACGGTGCT ATGGCTACAGGTTGGGCTAAAGTCAACGGTTCATGGTACTACCTAAACGCTAACGGTTCAATGGCAACAGGTTGGGTGAAAGA $\tt TGGAGATACCTGGTACTATCTTGAAGCATCAGGTGCTATGAAAGCCAATGGTTCAAAGTATCAGATAAATGGTACTATG$ TCAATGGCTTAGGTGCCCTTGCAGTCAACACACTGTAGATGGCTATAAAGTCAATGCCAATGGTGAATGGGTTTAA

(SEO, ID, NO, 184)

MNKKKMILTSLASVAILGAGFVTSQPTFVRAEESPQVVEKSSLEKKYEEAKAKADTAKKDYETAKKKAEDAQKKYEDDQKRTE
EKARKEAEASQKLNDVALVVQNAYKEYREVQNQRSKYKSDAEYQKKLTEVDSKIEKARKEQQDLQNKFNEVRAVVVPEPNALA
ETKKKAEEAKAEEKVAKRKYDYATLKVALAKKEVEAKELEIEKLQYEISTLEQEVATAQHQVDNLKKLLAGADPDDGTEVIEA
KLKKGEAELNAKQAELAKKQTELEKLLDSLDPEGKTQDELDKEAEEAELDKKADELQNKVADLEKEISNLEILLGGADPEDDT
AALQNKLAAKKAELAKKQTELEKLLDSLDPEGKTQDELDKEAEEAELDKKADELQNKVADLEKEISNLEILLGGADSEDDTAA
LQNKLATKKAELEKTQKELDAALNELGPDGDEEETPAPAPQPEQPAPAPKPEQPAPAPKPEQPAPAPKPEQPAPAPKPEQPAPAPKPEQPAPAPKPEQPAPAPKPEQPAPAPKPEQPAPAPKPEQPAPAPKPEQPAPAPKPEQPAPAPKPEQPAPAPKPEQPAPAPKPEQPAPATPKTGWKQENGMWYFYNTDGSMAIGWLQNNGSWYYLNANGAMATGWVKDGDTWYYLEAS
GAMKASQWFKVSDKWYYUNSNGAMATGWLQYNGSWYYLNANGDMATGWLQYNGSWYYLNANGAMATGWLQYN

ID302 (SEQ. ID. NO. 197)

ATGTTTGCATCAAAAAGCGAAAGAAAAGTACATTATTCAATTCGTAAATTTAGTGTTGGAGTAGCTAGTGTAGTTGTTGCCAG GTCAGGCAGAACAAGGAGAACAACCTAAAAAACTCGATTCAGAACGAGATAAGGCAAGGAAAGAGGTCGAGGAATATGTAAAA AAAATAGTGGGTGAGAGCTATGCAAAAATCAACTAAAAAGCGACATACAATTACTGTAGCTCTAGTTAACGAGTTGAACAACAT TAAGAACGAGTATTTGAATAAAATAGTTGAATCAACCTCAGAAAGCCAACTACAGATACTGATGATGGAGAGTCGATCAAAAG TAGATGAAGCTGTGTCTAAGTTTGAAAAGGACTCATCTTCTTCGTCAAGTTCAGACTCTTCCACTAAACCGGAAGCTTCAGAT ACAGCGAAGCCAAACAAGCCGACAGAACCAGGAGAAAAGGTAGCAGAAGCTAAGAAGAAGGTTGAAGAAGCTGAGAAAAAAGC ACGAAGAGCAGATGCTAAAGAGCAAGGTAAACCAAAGGGGCGGCAAAACGAGGAGTTCCTGGAGAGCTAGCAACACCTGATA AAAAAGAAAATGATGCGAAGTCTTCAGATTCTAGCGTAGGTGAAGAAACTCTTCCAAGCCCATCCCTGAAACCAGAAAAAAAG GTAGCAGAAGCTGAGAAGAAGGTTGAAGAAGCTAAGAAAAAAGCCGAGGATCAAAAAGAAGAAGATCGCCGTAACTACCCAAC CAATACTTACAAAACGCTTGAACTTGAAATTGCTGAGTCCGATGTGGAAGTTAAAAAAGCGGAGCTTGAACTAGTAAAAGAGG GAAAAAATCAAGACAGATCGTAAAAAAGCAGAAGAAGAAGCTAAACGAAAAGCAGCAGAAGAAGATAAAGTTAAAGAAAAACC AGCTGAACAACCACCAGCGCCCGGCTCCAAAAGCAGAAAAACCAGCTCCAGACAAACCAGAGAATCCAGCTGAACAAC CAGCAACCGCCAAAAACTGAAAAACCAGCACAACCATCTACTCCAAAAACAGGCTGGAAACAAGAAAACGGTATGTGGTACTT CTACAATACTGATGGTTCAATGGCGACAGGATGGCTCCAAAACAATGGCTCATGGTACTACCTCAACAGCAATGGCGCTATGG CGACAGGATGGCTCCAAAACAATGGTTCATGGTACTATCTAAACGCTAATGGTTCAATGGCAACAGGATGGCTCCAAAACAAT ${\tt GGTTCATGGTACTACCTAAACGCTAATGGTTCAATGGCGACAGGATGGCTCCAATACAATGGCTCATGGTACTAAACGC}$ ${\tt TAATGGTTCAATGGCGACAGGATGGCTCAATACAATGGCTCATGGTACTACCTAAACGCTAATGGTGATATGGCGACAGGTT}$ TGGTACTATGTCAATGGCTCAGGTGCCCTTGCAGTCAACACACTGTAGATGGCTATGGAGTCAATGCCAATGGTGAATGGGT AAACTAA

(SEO. ID. NO. 185)

MFASKSERKVHYSIRKFSVGVASVVVASLVMGSVVHATENEGATQVPTSSNRANESQAEQGEQPKKLDSERDKARKEVEEYVK KIVGESYAKSTKKRHTITVALVNELNNIKNEYLNKIVESTSESQLQILMMESRSKVDEAVSKFEKDSSSSSSSDSSTKPEASD TAKPNKPTEPGEKVAEAKKKVEEAEKKAKDQKEEDRRNYPTITYKTLELEIAESDVEVKKAELELVKVKANEPRDEQKIKQAE AEVESKQAEATRLKKIKTDREEAEEEAKRRADAKEQGKPKGRAKRGVPGELATPDKKENDAKSDSSVGEETLPSPSLKPEKK VAEAEKKVEEAKKKAEDQKEEDRRNYPTNTYKTLELEIAESDVEVKKAELELVKEEAKEPRNEEKVKQAKAEVESKKAEATRL EKIKTDRKKAEEEAKRKAAEEDKVKEKPAEQPQPAPAPKAEKPAPAPKPENPAEQPKAEKPADQQAEEDYARRSEEEYNRLTQ





QQPPKTEKPAQPSTPKTGWKQENGMWYFYNTDGSMATGWLQNNGSWYYLNSNGAMATGWLQNNGSWYYLNANGSMATGWLQNN GSWYYLNANGSMATGWLQYNGSWYYLNANGSMATGWLQYNGSWYYLNANGDMATGWVKDGDTWYYLEASGAMKASQWFKVSDK WYYVNGSGALAVNTTVDGYGVNANGEWVNZ

ID303 (SEO. ID. NO. 198)

 ${\tt ATGGTAAAAAGACGTATAAGGAGAGGGACGAGAAACCTGAAAAAGTTGTTGTTCCTGAGCAATCATCTATTCCTTCGTATCC}$ TGTATCTGTTACATCTAACCAAGGAACAGATGTAGCAGTAGAACCAGCTAAAGCAGTTGCTCCAACAACAGACTGGAAACAAG AAAATGGTATGTGGTATTTTTATAATACTGATGGTTCCATGGCAACAGGTTGGGTACAAGTTAATAGTTCATGGTACTACCTC AACAGCAACGGTTCTATGAAAGTCAATCAATGGTTCCAAGTTGGTGGTAAATGGTATTATGTAAATACATCGGGTGAGTTAGC GGTCAATACAAGTATAGATGGCTATAGAGTCAATGATAATGGTGAATGGGTGCGTTAA

MVKRRIRRGTREPEKVVVPEQSSIPSYPVSVTSNQGTDVAVEPAKAVAPTTDWKQENGMWYFYNTDGSMATGWVQVNSSWYYL NSNGSMKVNQWFQVGGKWYYVNTSGELAVNTSIDGYRVNDNGEWVRZ

ID304 (SEQ. ID. NO. 199)

 $\tt CTGAATACAAGTTTTGTTCATGCTGCTGATGGGATTCAATATGTCAGAGATGATACTAGAGATAAAGAAGAGGGAATAGAGTA$ $\tt TGATGACGCTGACAATGGGGATATTATTGTAAAAGTAGCGACTAAACCTAAGGTAGTAACCAAGAAAATTTCAAGTACGCGAA$ ${ t TTCGTTATGAAAAGATGAAAACAAAAGACCGTAGTGAAAAATCCTGTTACAATTGATGGAGAGGATGGCTATGTAACTACGACA$ ${f AGGACCTACGATGTTAATCCAGAGACTGGTTATGTTACCGAACAGGTTACTGTTGATAGAAAAGAAGCCACGGATACAGTTAT}$ CAAAGTTCCAGCTAAAAGCAAGGTTGAAGAAGTTCTTGTTCCATTTGCTACTAAATATGAAGCAGACAATGACCTTTCTGCAG GACAGGAGCAAGAGATTACTCTAGGAAAGAATGGGAAAACAGTTACAACGATAACTTATAATGTAGATGGAAAGAGTGGACAA AGAAATTCCAATCGAAACAGAATATCTCGATGGCCCAACTCTTGATAAAAGTCAAGAAGTAGAAGAAGTAGGAGAAATTGGTA AATTACTCTTACTACAATCTATACTGTAG

LNTSFVHAADGIQYVRDDTRDKEEGIEYDDADNGDIIVKVATKPKVVTKKISSTRIRYEKDETKDRSENPVTIDGEDGYVTTT VTESTLSQKKDSQTRVVKKRTKPQVLVQEIPIETEYLDGPTLDKSQEVEEVGEIGKLLLLQSILZ

ID305 (SEO. ID. NO. 200)
ATGAAGCTTTTGAAAAAAATGATGCAAATCGCACTAGCCACATTTTTCTTCGGTTTGTTAGCGACAAATACAGTATTTGCAGA TGATTCTGAAGGATGGCAGTTTGTCCAAGAAAATGGTAGAACCTACTACAAAAAGGGGGGATCTAAAAGAAACCTACTGGAGAG TGATAGATGGGAAGTACTATTATTTTGATCCTTTATCCGGAGAGATGGTTGTCGGCTGGCAATATATACCTGCTCCACACAAG ${\tt CAGAGAAACGAGTCTATTATTTTGAAGATCAGCGTAGTTATCATACTTTAAAAACTGGTTGGATTTATGAAGAGGGTCATTGG}$ TATTATTTACAGAAGGATGGTGGCTTTGATTCGCGCATCAACAGATTGACGGTTGGAGAGCTAGCACGTGGTTGGGTTAAGGA TTACCCTCTTACGTATGATGAAGAGCAACCTAAAAGCAGCTCCATGGTACTATCTAAATCCAGCAACTGGCATTATGCAAACAG AGGAGCTATGGCAACTGGTTGGTATCAGGAAAGTTCGACTTGGTACTATCTAAATGCAAGTAATGGAGATATGAAAACAGGCT GGTTCCAAGTCAATGGTAACTGGTACTATGCCTATGATTCAGGTGCTTTAGCTGTTAATACCACAGTAGGTGGTTACTACTTA AACTATAATGGTGAATGGGTTAAGTAA

(SEQ. ID. NO. 188)

MKLLKKMMQIALATFFFGLLATNTVFADDSEGWQFVQENGRTYYKKGDLKETYWRVIDGKYYYFDPLSGEMVVGWQYIPAPHK GVTIGPSPRIEIALRPDWFYFGQDGVLQEFVGKQVLEAKTATNTNKHHGEEYDSQAEKRVYYFEDQRSYHTLKTGWIYEEGHW YYLQKDGGFDSRINRLTVGELARGWVKDYPLTYDEEKLKAAPWYYLNPATGIMQTGWQYLGNRWYYLHSSGAMATGWYKEGST WYYLDAENGDMRTGWQNLGNKWYYLRSSGAMATGWYQESSTWYYLNASNGDMKTGWFQVNGNWYYAYDSGALAVNTTVGGYYL NYNGEWVKZ

ID306 (SEQ. ID. NO. 201)

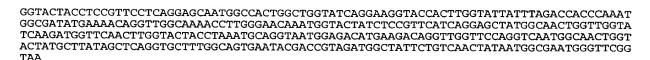
TTGGCTGGTAGATATGGTTCTGCTGTTCAGTGTACAGAAGTGACTGCCTCAAACCTTTCAACAGTTAAAACTAAAGCTACGGT TGTAGAAAAACCACTGAAAGATTTTAGAGCGTCTACGTCTGATCAGTCTGGTTGGGTGGAATCTAATGGTAAATGGTATTTCT ATGAGTCTGGTGATGTGAAGACAGGTTGGGTGAAAACAGATGGTAAATGGTACTATTTGAATGACTTAGGTGTCATGCAGACT GTATCATGAAGACAGGTTGGTTTAAAGTCGGACCACACTGGTACTATGCCTACGGTTCAGGAGCTTTGGCTGTGAGCACAACA ACACCAGATGGTTACCGTGTAAATGGTAATGGTGAATGGGTAAACTAG

(SEQ. ID. NO. 189)

LAGRYGSAVQCTEVTASNLSTVKTKATVVEKPLKDFRASTSDQSGWVESNGKWYFYESGDVKTGWVKTDGKWYYLNDLGVMQT GFVKFSGSWYYLSNSGAMFTGWGTDGSRWFYFDGSGAMKTGWYKENGTWYYLDEAGIMKTGWFKVGPHWYYAYGSGALAVSTT TPDGYRVNGNGEWVNZ

ID307 (SEQ. ID. NO. 202)

 ${f ATGAAAATTTTGAAAAAAACTATGCAAGTTGGACTGACAGTATTTTTCTTTGGTTTGCTAGGGACCAGTACAGTATTTGCAGA$ TGATTCTGAAGGATGGCAGTTTGTCCAAGAAAACGGAAGAACCTACTACAAAAAGGGGGGACCTCAAAGAAACCTACTGGCGAG ${\tt AGTGCTACAAGAGTTTGTTGGTTGGAAAACATTAGAGATTAAAACTAAAGACAGTGTTGGAAGAAAGTACGGGGAAAAACGTG}$ AAGATTCAGAAGATAAAGAAGAGAAGCGTTATTATACGAACTATTACTTTAATCAAAATCATTCTTTAGAGACAGGTTGGCTT TATGATCAGTCTAACTGGTATTATCTAGCTAAGACGGAAATTAATGGAGAAAACTACCTTGGTGGTGAAAGACGTGCGGGGTG GATAAACGATGATTCGACTTGGTACTACCTAGATCCAACAACTGGTATTATGCAAACAGGTTGGCAATATCTAGGTAATAAGT



(SEQ. ID. NO. 190)

MKILKKTMQVGLTVFFFGLLGTSTVFADDSEGWQFVQENGRTYYKKGDLKETYWRVIDGKYYYFDSLSGEMVVGWQYIPFPSK GSTIGPYPNGIRLEGFPKSEWYYFDKNGVLQEFVGWKTLEIKTKDSVGRKYGEKREDSEDKEEKRYYTNYYFNQNHSLETGWL YDQSNWYYLAKTEINGENYLGGERRAGWINDDSTWYYLDPTTGIMQTGWQYLGNKWYYLRSSGAMATGWYQEGTTWYYLDHPN GDMKTGWQNLGNKWYYLRSSGAMATGWYQDGSTWYYLNAGNGDMKTGWFQVNGNWYYAYSSGALAVNTTVDGYSVNYNGEWVR

ID308 (SEO, ID. NO. 203)

 $\overline{\textbf{ATGAAAAAGAAATTAACT}} \textbf{AGTTTAGCACTTGTAGGCGCTTTTTTAGGTTTGTCATGGTATGGGAATGTTCAGGCTCAAGAAAG}$ ${ t TTCAGGAAATAAAATCCACTTTATCAATGTTCAAGAAGGTGGCAGTGATGCGATTATTCTTGAAAGCAATGGACATTTTGCCA$ TGGTGGATACAGGAGAAGATTATGATTTCCCAGATGGAAGTGATTCTCGCTATCCATGGAGAGAAGGAATTGAAACGTCTTAT AAGCATGTTCTAACAGACCGTGTCTTTCGTCGTTTGAAGGAATTGGGTGTCCAAAAACTTGATTTTATTTTGGTGACCCATAC ${\tt CCACAGTGATCATATTGGAAATGTTGATGAATTACTGTCTACCTATCCAGTTGACCGAGTCTATCTTAAGAAATATAGTGATA}$ GTCGTATTACTAATTCTGAACGTCTATGGGATAATCTGTATGGCTATGATAAGGTTTTACAGACTGCTGCAGAAAAAGGTGTT TCAGTTATTCAAAATATCACACAAGGGGATGCTCATTTTCAGTTTGGGGACATGGATATTCAGCTCTATAATTATGAAAATGA AACTGATTCATCGGGTGAATTAAAGAAAATTTGGGATGACAATTCCAATTCCTTGATTAGCGTGGTGAAAGTCAATGGCAAGA AAGTTTAATCATCACCATGATACCAACAAATCAAATACCAAGGATTTCATTAAAAATTTGAGTCCGAGTTTGATTGTTCAAAC TTCGGATAGTCTACCTTGGAAAAATGGTGTTGATAGTGAGTATGTTAATTGGCTCAAAGAACGAGGAATTGAGAGAATCAACG CCAAGTTTTCAAGCTGGTTGGCATAAGAGTGCATATGGGAACTGGTGGTATCAAGCGCCTGATTCTACAGGAGAGTATGCTGT CGGTTGGAATGAAATCGAAGGTGAATGGTATTACTTTAACCAAACGGGTATCTTGTTACAGAATCAATGGAAAAAATGGAACA ATCATTGGTTCTATTTGACAGACTCTGGTGCTTCTGCTAAAAATTGGAAGAAAATCGCTGGAATCTGGTATTATTTTAACAAA $\hbox{\tt GAAAACCAGATGGAAATTGGTTGGATTCAAGATAAAGAGCAGTGGTATTATTTGGATGTTGATGGTTCTATGAAGACAGGATG}$ GCTTCAATATATGGGGCAATGGTATTACTTTGCTCCATCAGGGGAAATGAAAATGGGCTGGGTAAAAGATAAAGAAACCTGGT ACTATATGGATTCTACTGGTGTCATGAAGACAGGTGAGATAGAAGTTGCTGGTCAACATTATTATCTGGAAGATTCAGGAGCT $\tt GTGGTGCCTGGTTAGTGGATGTTTCGATCGAGAAATCTGCTACAATTAAAACTACAAGTCATTCAGAAATAAAAGAATCCAAA$ GAAGTAGTGAAAAAGGATCTTGAAAATAAAGAAACGAGTCAACATGAAAGTGTTACAAATTTTTCAACTAGTCAAGATTTGAC ATCCTCAACTTCACAAAGCTCTGAAACGAGTGTAAACAAATCGGAATCAGAACAGTAG

(SEQ. ID. NO. 191)

MKKKLTSLALVGAFLGLSWYGNVQAQESSGNKIHFINVQEGGSDAIILESNGHFAMVDTGEDYDFPDGSDSRYPWREGIETSY KHVLTDRVFRRLKELGVQKLDFILVTHTHSDHIGNVDELLSTYPVDRVYLKKYSDSRITNSERLWDNLYGYDKVLQTAAEKGV SVIQNITQGDAHFQFGDMDIQLYNYENETDSSGELKKIWDDNSNSLISVVKVNGKKIYLGGDLDNVHGAEDKYGPLIGKVDLM KFNHHHDTNKSNTKDFIKNLSPSLIVQTSDSLPWKNGVDSEYVNWLKERGIERINAASKDYDATVFDIRKDGFVNISTSYKPI PSFQAGWHKSAYGNWWYQAPDSTGEYAVGWNEIEGEWYYFNQTGILLQNQWKKWNNHWFYLTDSGASAKNWKKIAGIHYYFNK ENQMEIGWIQDKEQWYYLDVDGSMKTGWLQYMGQWYYFAPSGEMKMGWVKDKETWYYMDSTGVMKTGEIEVAGQHYYLEDSGA MKQGWHKKANDWYFYKTDGSRAVGWIKDKDKWYFLKENGQLLVNGKTPEGYTVDSSGAWLVDVSIEKSATIKTTSHSEIKESK EVVKKDLENKETSQHESVTNFSTSQDLTSSTSQSSETSVNKSESEQZ

ID309 (SEQ. ID. NO. 204)

ATGGAAATTAATGTGAGTAAATTAAGAACAGATTTGCCTCAAGTCGGCGTGCAACCATATAGGCAAGTACACGCACACTCAAC
TGGGAATCCGCATTCAACCGTACAGAATGAAGCGGATTATCACTGGCGGAAAGACCCAGAATTAGGTTTTTTCTCGCACATTG
TTGGGAACCGTTTCAACCGTACAGAATGAAGCCGGTTGATAATGGTGCCTGGGACCGTTGGGACCGTTTTTTTCTCTGCACATTG
CCAGCGGTTGAACTGATTGAAAGCCATTCAACCAAAGAAGAGTTCATGACGGACTACCGCCTTTATATCGAACTTTCACGCAA
TCTAGCAGATGAAGCAGGTTTGCCGAAAACCGCTTGATACAGGGAGTTTAGCCGGATTAAAACGCACGAGTATTGCACGAATA
ACCAAACAAACAACACCACCTCCAGACCACCTTGACCCTTATCCATATCTTGCTAAATGGGGCATTAGCCGTGAGCAGTTTAAGCAT
GATATTGAGAACAGCCTCGACACCGCTTGACCCTTATCCATATCTTGCTAAATGGGCTACCTTCAGACGCTCTTA
TCCAAAAGACAACGTTGACGATTGAAACAGGCTGGCAGAAGAATGACACTGGCTACTTGCAGACCACTCGGAGAAGAAAATCACTCAGACGACTTTGGCACACTTTGACACTTTTGACAGTTCAGGCTATATGCTTGCAGACCGCTGGAGGA
AGCACACAGACGGCAACTGGTACTGGTTCGACACACTCAGGCGAAATGGCTACAGGCTGGAAGAAAATCGCTGATAAGTGGTAC
TATTTCAACGAAGAAGGTGCCATGAAGACAGGCTGGGTCAAAGTACAAGGACACTTGGTACTACTTAGACGCTAAAGAAGACGCC
CATGGTATCAAATGCCTTTATCCAGTCAGCGGACGGACAGGCTGGTACTACCTCAAACCAGACGGAACACTGGCAGACAACC
CAGAATTCACAGTAGAGCCAGATGGCTTGATTACAGTAAAATAA

(SEQ. ID. NO. 192)

MEINVSKLRTDLPQVGVQPYRQVHAHSTGNPHSTVQNEADYHWRKDPELGFFSHIVGNGCIMQVGPVDNGAWDVGGGWNAETY AAVELIESHSTKEEFMTDYRLYIELLRNLADEAGLPKTLDTGSLAGIKTHEYCTNNQPNNHSDHVDPYPYLAKWGISREQFKH DIENGLTIETGWQKNDTGYWYVHSDGSYPKDKFEKINGTWYYFDSSGYMLADRWRKHTDGNWYWFDNSGEMATGWKKIADKWY YFNEEGAMKTGWVKYKDTWYYLDAKEGAMVSNAFIQSADGTGWYYLKPDGTLADKPEFTVEPDGLITVKZ

ID310 (SEQ. ID. NO. 205)



(SEQ. ID. NO. 193)

MGTTGFTIIDLIILIVYLLAVLVAGIYFSKKEMKGKEFFKGDGSVPWYVTSVSIFATMLSPISFLGLAGSSYAGSWILWFAQL
GMVVAIPLTIRFILPIFARIDIDTAYDYLDKRFNSKALRIISALLFIIYQLGRMSIIMYLPSAGLSVLTGIDINILIILMGVV
AIVYSYTGGLKSVLWTDFIQGVILISGVVLALFVLIANIKGGFGAVAETLANGKFLAANEKLFDPNLLSNSIFLIVMGSGFTI
LSSYASSQDLVQRFTTTQNIKKLNKMLFTNGVLSLATATVFYLIGTGLYVFYQVQNADSAASNIPQDQIFMYFIAYQLPVGIT
GLILAAIYAASQSTISTGLNSVATSWTLDIQDVISKNMSDNRRTKIAQFVSLAVGLFSIGVSIVMAHSDIKSAYEWFNSFMGL
VLGLLGGVFILGFVSKKANKQGAYAALIVSTIVMVFIKYFLPPTAVSYWAYSLISISVSVVSGYIVSVLTGNKVSAPKYTTIH
DITEIKADSSWEVRHZ

ID311 (SEQ. ID. NO. 206)

AGCTGGTCAGGATAAGAAAGAGTCTAATCGAGTTGCTTATATAGATGGTGATCAGGCTGGTCAAAAGGCAGAAAACTTGACAC ${\tt CAGATGAAGTCAGTAAGAGGGGAGGGGATCAACGCCGAACAAATCGTCATCAAGATTACGGATCAAGGTTATGTGACCTCTCAT}$ GGAGACCATTATCATTACTATAATGGCAAGGTCCCTTATGATGCCATCATCAGTGAAGAGCTCCTCATGAAAGATCCGAATTA TCAGTTGAAGGATTCAGACATTGTCAATGAAATCAAGGGTGGTTATGTCATCAAGGTAGACGGAAAATACTATGTTTACCTTA AGGATGCAGCTCATGCGGATAATATTCGGACAAAAGAAGAGATTAAACGTCAGAAGCAGGAACGCAGTCATAATCACGGGTCA GGAGCTAACGATCATGCAGTAGCTGCAGCCAGAGCCCAAGGACGCTATACAACGGATGATGGGTATATCTTCAATGCATCTGA TATCATTGAGGACACGGGTGATGCTTATATCGTTCCTCACGGCGACCATTACCATTACATTCCTAAGAATGAGTTATCAGCTA GCGAGTTAGCTGCTGCAGAAGCCTATTGGAATGGGAAGCAGGGATCTCGTCCTTCTTCAAGTTCTAGTTATAATGCAAATCCA GCTCAACCAAGATTGTCAGAGAACCACAATCTGACTGTCACTCCAACTTATCATCAAAAATCAAGGGGAAAACATTTCAAGCCT ${\tt TTTACGTGAATTGTATGCTAAACCCTTATCAGAACGCCATGTGGAATCTGATGGCCTTATTTTCGACCCAGCGCAAATCACAA}$ GTCGAACCGCCAGAGGTGTAGCTGTCCCTCATGGTAACCATTACCACTTTATCCCTTATGAACAAATGTCTGAATTGGAAAAA ${\tt CGAATTGCTCGTATTATTCCCCTTCGTTATCGTTCAAACCATTGGGTACCAGATTCAAGACCAGAACAACCAAGTCCACAATC}$ GACTCCGGAACCTAGTCCAAGTCCGCAACCTGCACCAAATCCTCAACCAGCTCCAAGCAATCCAATTGATGAGAAATTGGTCA AAGAAGCTGTTCGAAAAGTAGGCGATGGTTATGTCTTTGAGGAGAATGGAGTTTCTCGTTATATCCCAGCCAAGGATCTTTCA GCAGAAACAGCAGCAGCATTGATAGCAAACTGGCCAAGCAGGAAAGTTTATCTCATAAGCTAGGAGCTAAGAAAACTGACCT CCCATCTAGTGATCGAGAATTTTACAATAAGGCTTATGACTTACTAGCAAGAATTCACCAAGATTTACTTGATAATAAAGGTC ${\tt GACAAGTTGATTTTGAGGCTTTGGATAACCTGTTGGAACGACTCAAGGATGTCCCAAGTGATAAAGTCAAGTTAGTGGATGAT}$ ATTCTTGCCTTCTTAGCTCCGATTCGTCATCCAGAACGTTTAGGAAAACCAAATGCGCAAATTACCTACACTGATGATGAGAT TCAAGTAGCCAAGTTGGCAGGCAAGTACACAACAGAAGACGGTTATATCTTTGATCCTCGTGATATAACCAGTGATGAGGGGG ATGCCTATGTAACTCCACATATGACCCATAGCCACTGGATTAAAAAAGATAGTTTGTCTGAAGCTGAGAGAGCGGCAGCCCAG GCTTATGCTAAAGAGAAAGGTTTGACCCCTCCTTCGACAGACCATCAGGATTCAGGAAATACTGAGGCAAAAGGAGCAGAAGC TTTTGGTAACGCTAGCGACCATGTTCAAAGAAACAAAAATGGTCAAGCTGATACCAATCAAACGGAAAAACCAAGCGAGGAGA AACCTCAGACAGAAAACCTGAGGAAGAAACCCCTCGAGAAGAGAGAACCGCAAAGCGAGAAACCAGAGTCTCCAAAACCAACA GAGGAACCAGAAGAATCACCAGAGGAATCAGAAGAACCTCAGGTCGAGACTGAAAAAGGTTGAAGAAAAACTGAGAGAGGCTGA AGATTTACTTGGAAAAATCCAGGATCCAATTATCAAGTCCAATGCCAAAGAGACTCTCACAGGATTAAAAAATAATTTACTAT

<u>(SEQ. ID. NO. 194)</u>

MKINKKYLAGSVAVLALSVCSYELGRHQAGQDKKESNRVAYIDGDQAGQKAENLTPDEVSKREGINAEQIVIKITDQGYVTSH
GDHYHYYNGKVPYDAIISEELLMKDPNYQLKDSDIVNEIKGGYVIKVDGKYYVYLKDAAHADNIRTKEEIKRQKQERSHNHGS
GANDHAVAAARAQGRYTTDDGYIFNASDIIEDTGDAYIVPHGDHYHYIPKNELSASELAAAEAYWNGKQGSRPSSSSSYNANP
AQPRLSENHNLTVTPTYHQNQGENISSLLRELYAKPLSERHVESDGLIFDPAQITSRTARGVAVPHGNHYHFIPYEQMSELEK
RIARIIPLRYRSNHWVPDSRPEQPSPQSTPEPSPSPQPAPNPQPAPSNPIDEKLVKEAVRKVGDGYVFEENGVSRYIPAKDLS
AETAAGIDSKLAKQESLSHKLGAKKTDLPSSDREFYNKAYDLLARIHQDLLDNKGRQVDFEALDNLLERLKDVPSDKVKLVDD
LLAFLAPIRHPERLGKPNAQITYTDDEIQVAKLAGKYTTEDGYIFDPRDITSDEGDAYVTPHMTHSHWIKKDSLSEAERAAAQ
AYAKEKGLTPPSTDHQDSGNTEAKGAEAIYNRVKAAKKVPLDRMPYNLQYTVEVKNGSLIIPHYDHYHNIKFEWFDEGLYEAP
KGYTLEDLLATVKYYVEHPNERPHSDNGFGNASDHVQRNKNGQADTNQTEKPSEEKPQTEKPEEETPREEKPQSEKPESPKPT
EEPEESPEESEEPQVETEKVEEKLREAEDLLGKIQDPIIKSNAKETLTGLKNNLLFGTQDNNTIMAEAEKLLALLKESKZ

ID312 (SEQ. ID. NO. 207)

(SEQ. ID. NO. 195)
MEGLVRVHLLPVFGDYKLSKLTTPILQQQVNKWADKANKGEKGAFANYSLLHNMNKRILKYGVAIQVIQYNPANDVIVPRKQQ
KEKAAVKYLDNKELKQFLDYLDALDQSNYENLFDVVLYKTLLATGCRISEALALEWSDIDLESGVISINKTLNRYQEINSPKS
SAGYRDIPIDKATLLLLKQYKNRQQIQSWKLGRSETVVFSVFTEKYAYACNLRKRLNKHFDAAGVTNVSFHGFRHTHTTMMLY
AQVSPKDVQYRLGHSNLMITENTYWHTNQENAKKAVSNYETAINNLZ

At Page 1, 2nd full paragraph:

Background of the Invention

Streptococcus pneumoniae, commonly referred to as the pneumococcus, is an important pathogenic organism. The continuing significance of Streptococcus pneumoniae infections in relation to human disease in developing and developed countries has been authoritatively reviewed (Fiber, G.R., Science, 265:1385-1387 (1994)). That indicates that on a global scale this organism is believed to be the most common bacterial cause of acute respiratory infections, and is estimated to result in 1 million childhood deaths each year, mostly in developing countries (Stansfield, S.K., Pediatr. Infect. Dis., 6:622 (1987)). In the USA it has been suggested (Breiman et al, Arch. Intern. Med., 150:1401 (1990)) that the pneumococcus is still the most common cause of bacterial pneumoniae, and that disease rates are particularly high in young children, in the elderly, and in patients with predisposing conditions such as asplenia, heart, lung, and kidney disease, diabetes, alcoholism, or with immunosuppressive disorders, especially AIDS. These groups are at higher risk of pneumococcal septicaemia and hence meningitis and therefore have a greater risk of dying from pneumococcal infection. The pneumococcus is also the leading cause of otitis media and sinusitis, which remain prevalent infections in children in developed countries, and which incur substantial costs.

At Page 3, 2nd full paragraph:

Brief Summary of the Invention

Thus, in a first aspect the present invention provides a Streptococcus pneumoniae protein or polypeptide having a sequence selected from those shown in Table 2 herein.

At Page 3, 2nd full paragraph:

Detailed Description of the Invention

As discussed herein, the proteins and polypeptides of the invention are useful as antigenic material. Such material can be "antigenic" and/or "immunogenic". Generally, "antigenic" is taken to mean that the protein or polypeptide is capable of being used to raise antibodies or indeed is capable of inducing an antibody response in a subject. "Immunogenic" is taken to mean that the protein or polypeptide is capable of eliciting a protective immune response in a subject. Thus, in the latter case, the protein or polypeptide may be capable of not only generating an antibody response and in addition non-antibody based immune responses.

At page 101, line 1:

CLAIMSWhat is claimed is: